

(21) Application No: **0722767.1**

(22) Date of Filing: **19.08.2004**

Date Lodged: **20.11.2007**

(30) Priority Data:  
(31) **10646682** (32) **22.08.2003** (33) **US**

(62) Divided from Application No  
**0418542.7** under Section 15(4) of the Patents Act 1977

(71) Applicant(s):  
**Dennis Sunga Fernandez**  
**1175 Osborn Avenue, Atherton,**  
**California 94027, United States of America**

(72) Inventor(s):  
**Dennis Sunga Fernandez**

(74) Agent and/or Address for Service:  
**Langner Parry**  
**High Holborn House, 52-54 High Holborn,**  
**LONDON, WC1V 6RR, United Kingdom**

(51) INT CL:  
**G06F 19/00** (2006.01) **A61B 5/00** (2006.01)  
**G01N 33/48** (2006.01)

(52) UK CL (Edition X ):  
**NOT CLASSIFIED**

(56) Documents Cited:  
**WO 2003/043684 A** **US 6178349 A**

(58) Field of Search:  
Other: **ONLINE: EPODOC, WPI**

(54) Abstract Title: **Systems biology based therapeutic modeling and implantable devices**

(57) A computer based method of predicting therapeutic strategy comprising representing one or more biological processes and molecular interactions as mathematical models and accessing a systems biology platform via a network to simulate drug effects and predict outcomes of interactions with the drug to aid target discovery and validation. Also claimed is an implantable network biosensor device comprising a sensor unit for receiving a multi-sensor signal from a biosensor platform detecting a biological material of a host, a controller for processing a systems-biology platform for simulation modelling of the biological material and a reservoir for releasing a therapeutic to the host wherein the sensor may automatically detect an effect of the therapeutic on the host.

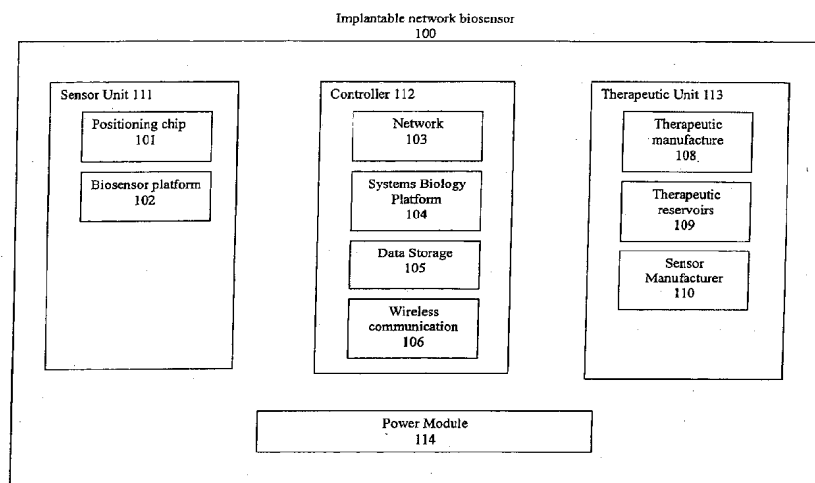


Figure 1a

Implantable network biosensor  
100

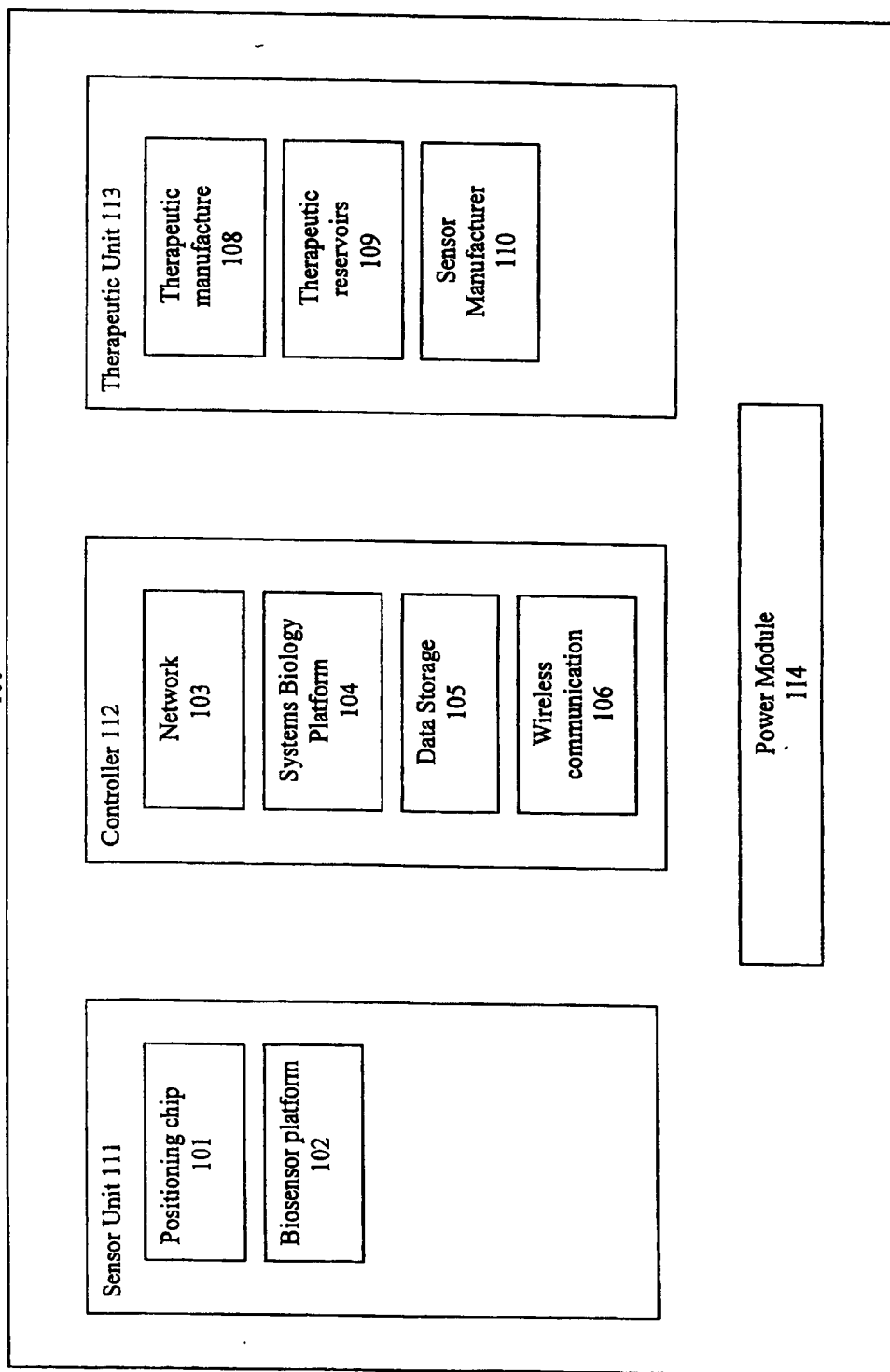


Figure 1a

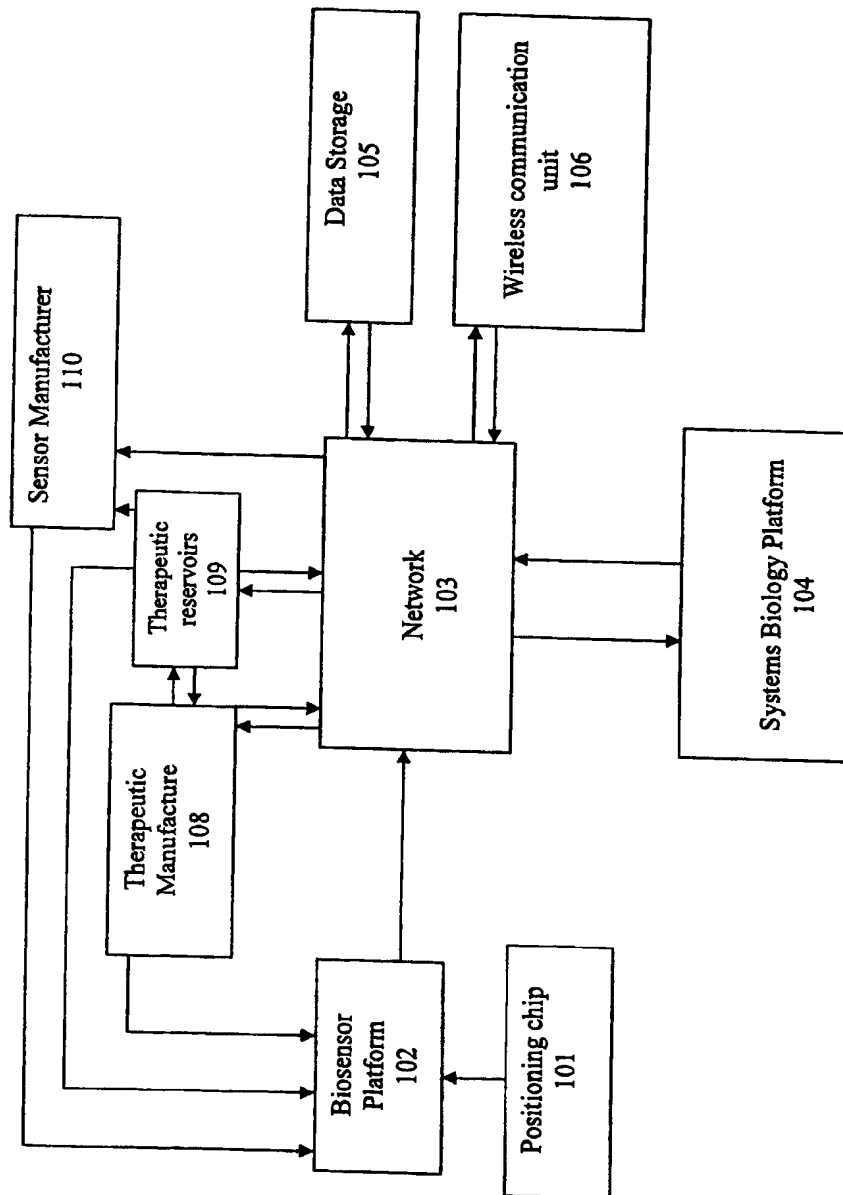
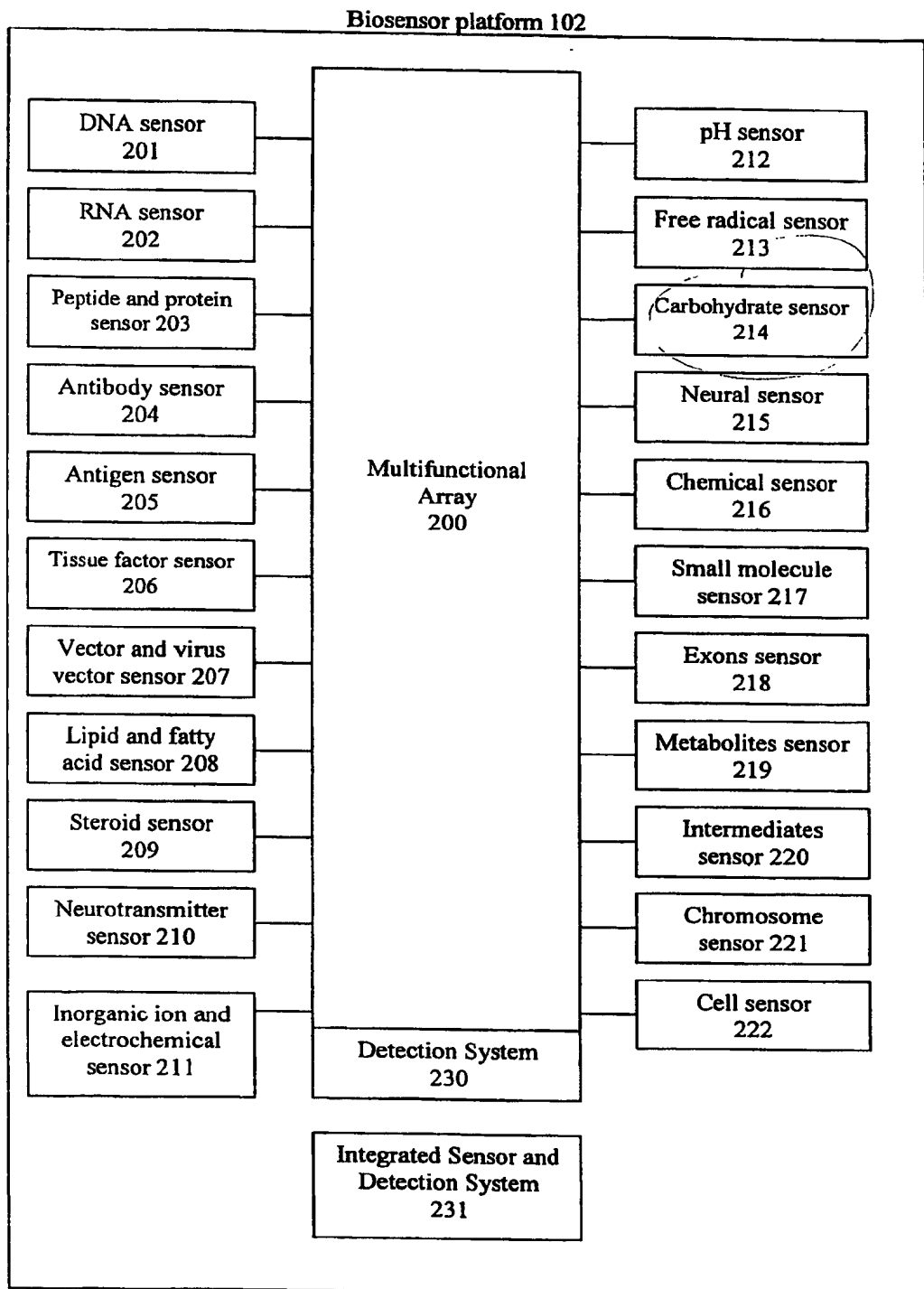
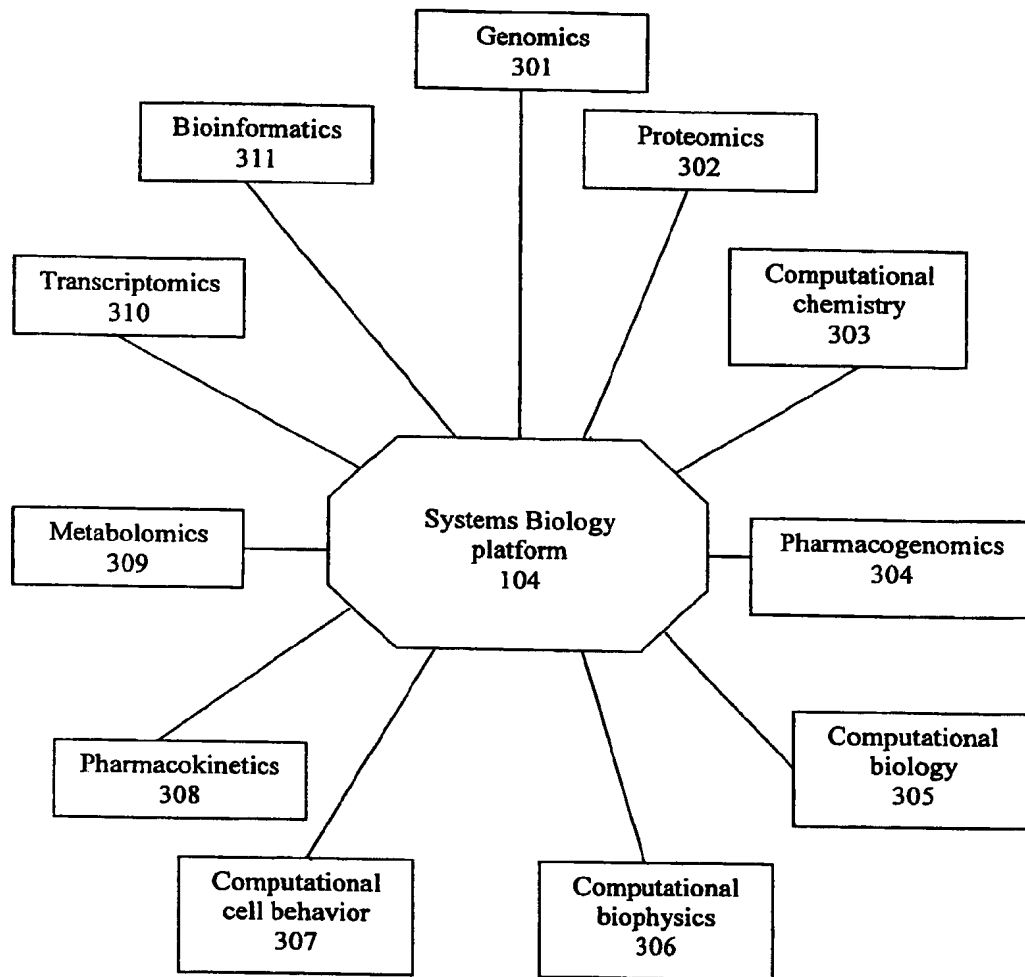
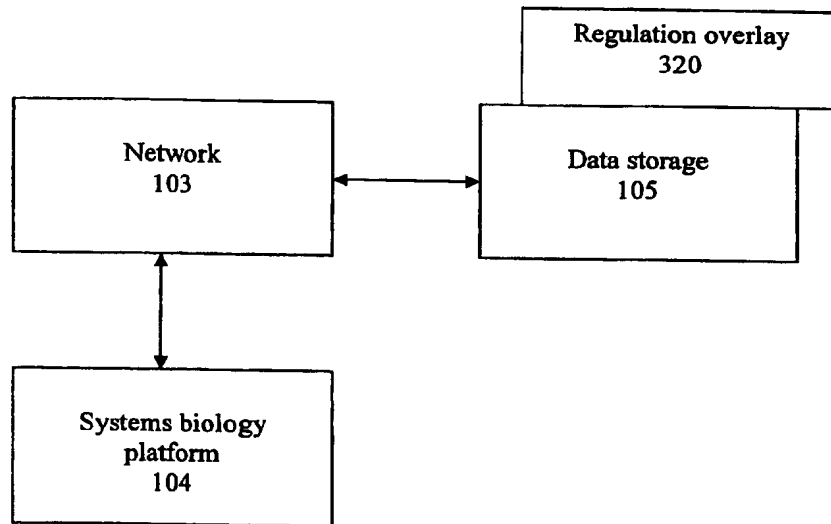


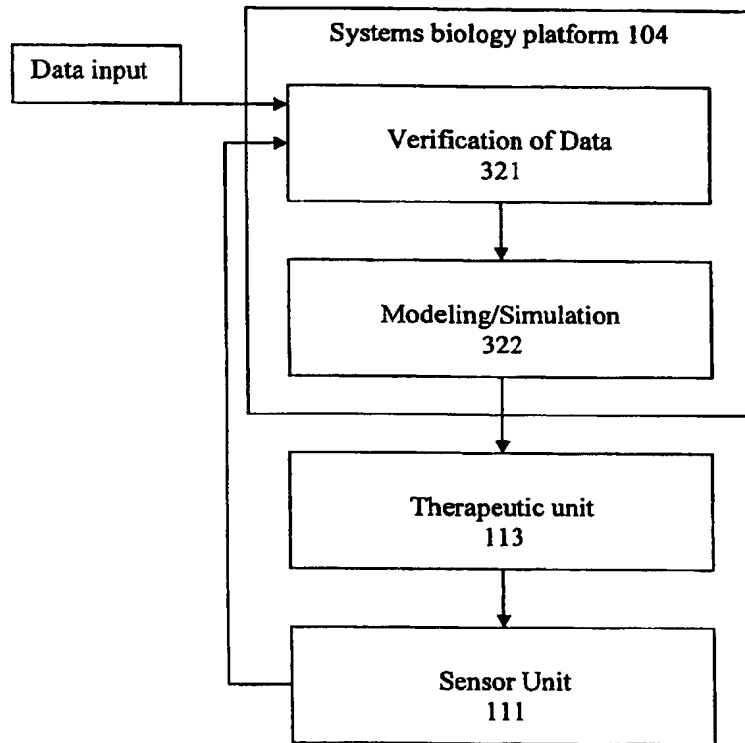
Figure 1b

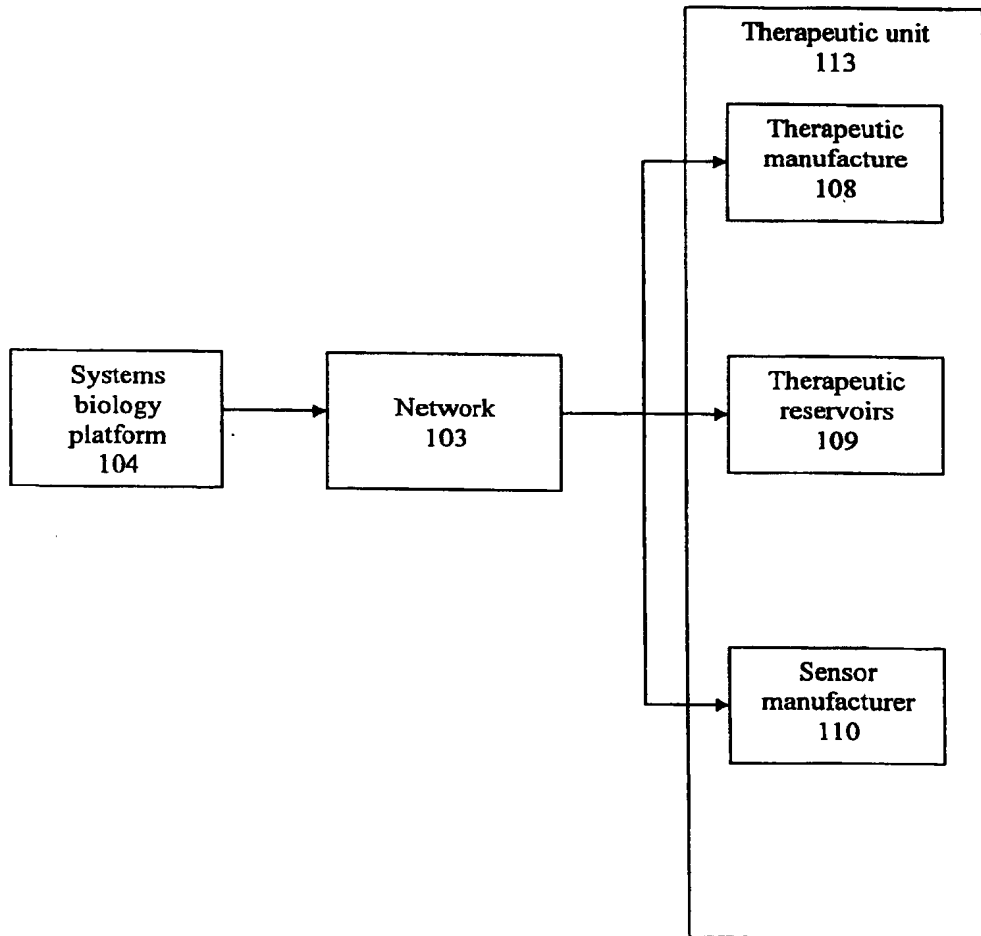
**Figure 2**

**Figure 3a**



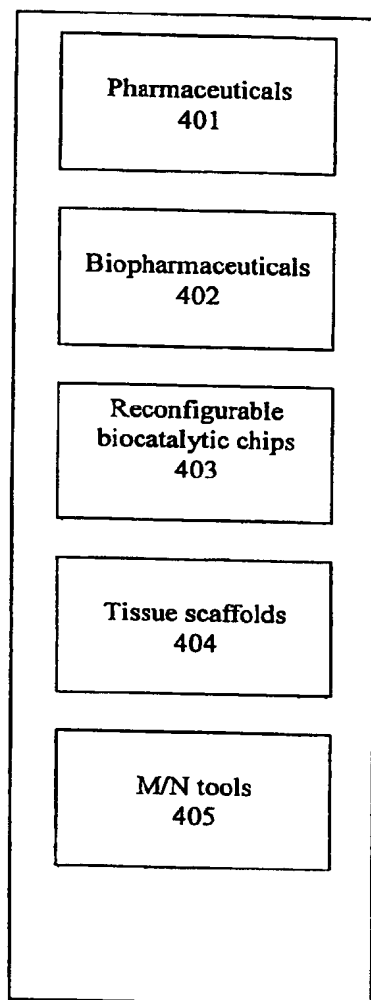
**Figure 3b**

**Figure 3c**

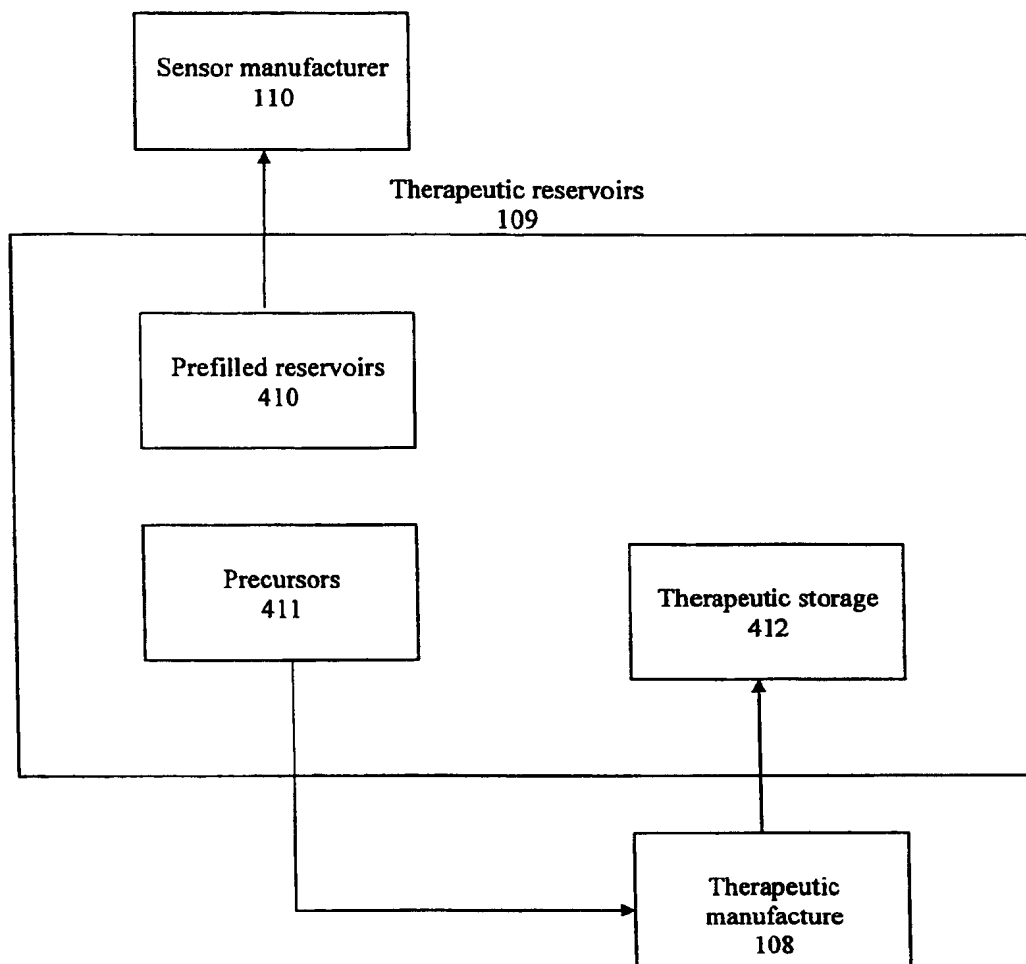
**Figure 4a**



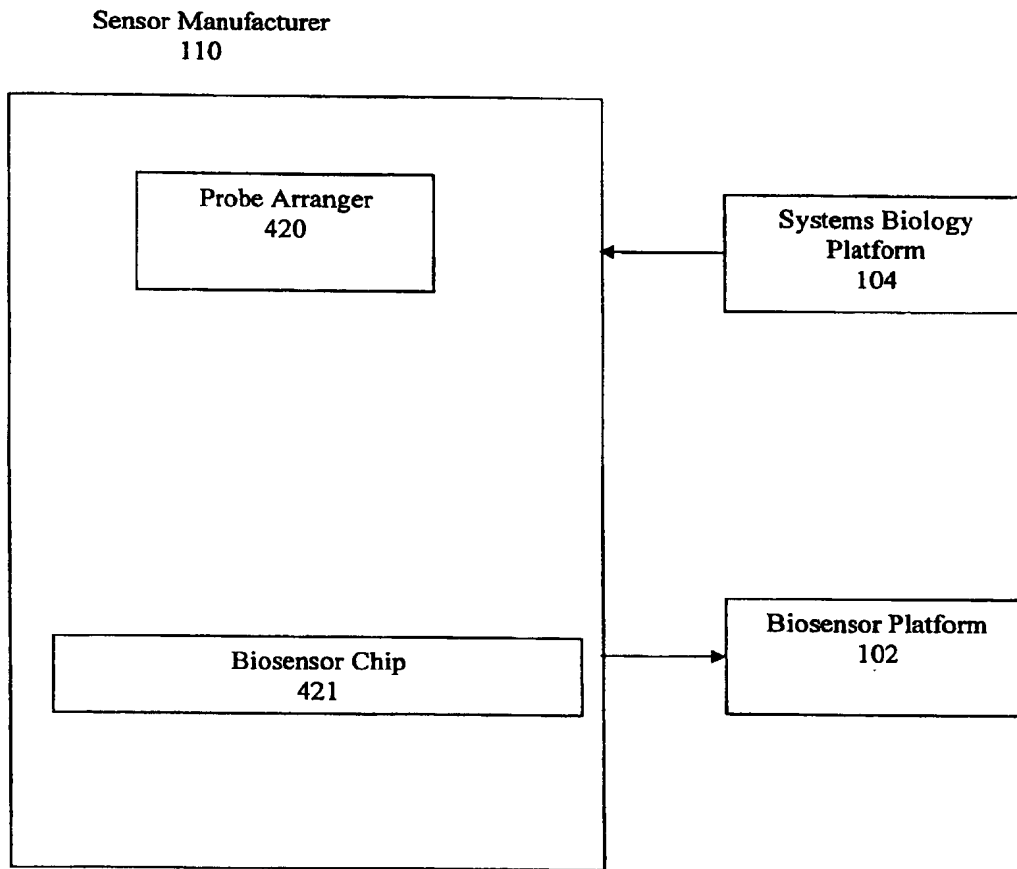
**Therapeutic Manufacture 108**



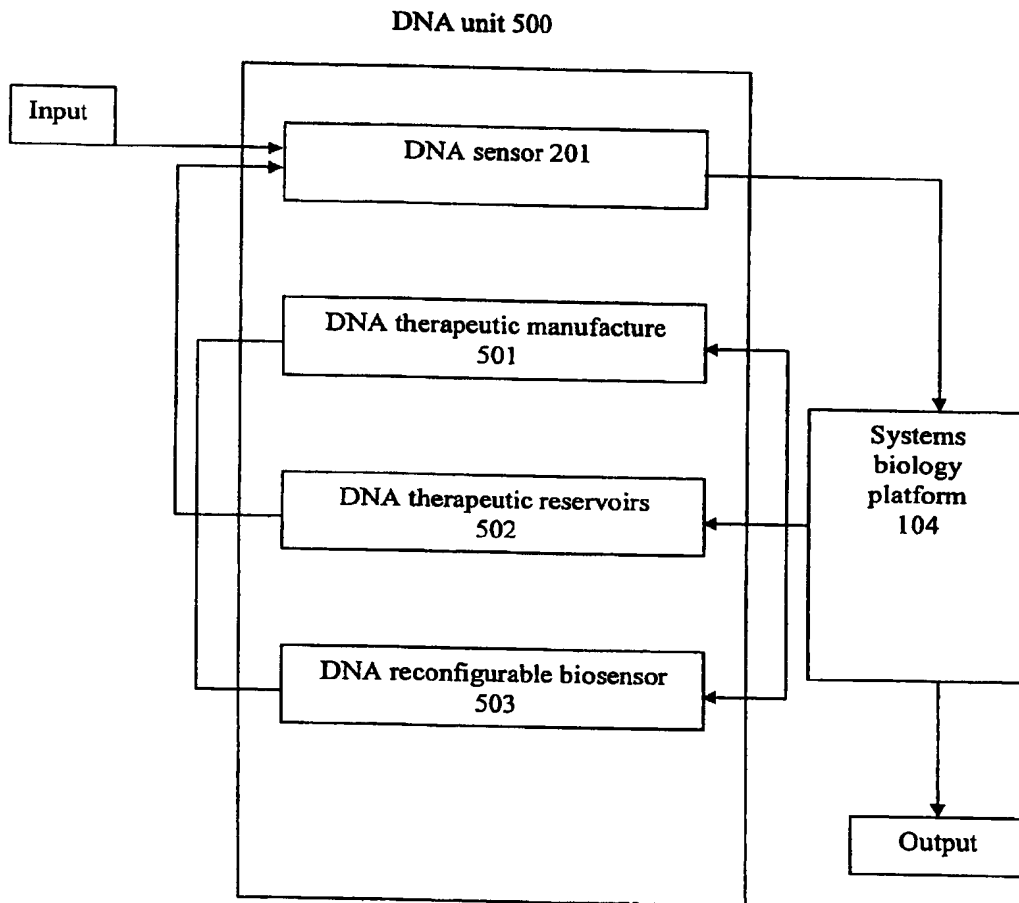
**Figure 4b**

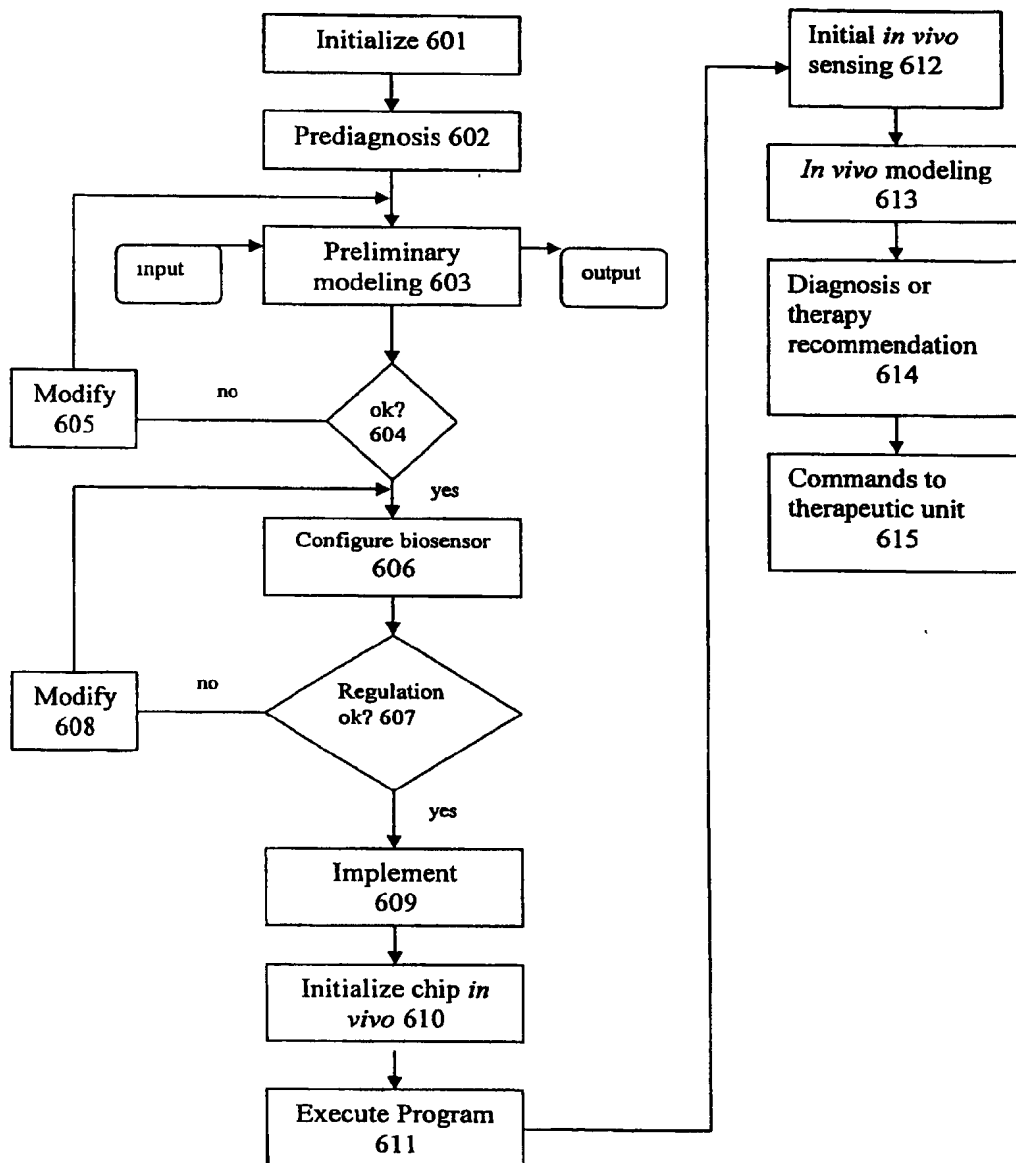


**Figure 4c**



**Figure 4d**

**Figure 5**



**Integrated Biosensor and Simulation System for Diagnosis and  
Therapy**

Invention relates to sensors for monitoring or analyzing  
5 biological hosts or material.

The present invention is divided from GB 0418542.8  
(published as GB 2405203 A) to which reference is directed.

10 Various sensors are used to detect or measure  
macroscopic or molecular physiology in humans or other  
biological host. Additionally systems-biology software  
provides computational modeling of molecular structures and  
interactions for genomics, proteomics, metabolomics,  
15 transcriptomics, computational chemistry, pharmacogenomics, or  
other purpose. Such tools, however, are not easily or  
automatically integrated or reconfigurable for  
interdisciplinary diagnosis or therapy.

20 Integrated biosensor-simulation system combines one or  
more sensor to detect various conditions in biological  
target or host, and software program or simulator using  
system-biology model and sensor data adaptively to provide  
therapy, diagnosis, or other automated feedback. Preferably  
25 one or more sensor is reconfigurable by the simulator.  
Optionally food material for consumption by the biological  
target is sensed for application to the simulator, which  
may apply certain regulatory condition. Switch couples  
simulator programmably to sensors.

30

Figure 1a shows sensor device according to aspect of  
present invention.

Figure 1b shows sensor network according to aspect of present invention.

5        Figure 2 shows sensor system according to aspect of present invention.

Figure 3a shows systems-biology software according to aspect of present invention.

10

Figure 3b shows systems biology software and data according to aspect of present invention.

Figure 3c shows system biology software and sensor  
15 according to aspect of present invention.

Figure 4a shows system biology software according to aspect of present invention.

20        Figure 4b shows therapy according to aspect of present invention.

Figure 4c shows therapy reservoir according to aspect of present invention.

25

Figure 4d shows sensor reconfiguration arrangement according to aspect of present invention.

Figure 5 shows DNA sensor according to aspect of  
30 present invention.

Figure 6 shows diagnosis or therapy method according to aspect of present invention.

Figure 1a architectural diagram illustrates implantable network biosensor 100. It is contemplated herein that sensor 100 may also operate without being implanted in biological host, but instead through external contact or attachment thereto. Optionally multiple coupled sensors 100 may provide fault-tolerant back-up or recovery facility, in case one or more sensors fails or malfunctions. Sensor 100 may be provided inside a host, e.g., mouth, larynx, blood vessel, vein, nose, ear, eye, heart, brain, lymph node, lung, breast, stomach, pancreas, kidney, colon, rectum, ovary, uterus, bladder, prostate, or other organ or using portable mobile application externally, e.g. skin, fingernail.

Sensor 100 includes sensor unit 111, controller 112, therapeutic unit 113, and power module 114. Sensor 100 components may be interconnected or communicate with other components using electrical, electronic, or electromagnetic signals, e.g., optical, radio frequency, digital, analog or other signalling scheme. Power module 114 provides electrical energy for sensor 100 to operate.

Generally biosensor 100 may sense individual genome, proteome, metabolism, transcription, translation, blood pressure, carbohydrate and oxygen concentrations, or other factors as described herein. Data is provided by sensor 100 to integrated network 103 that applies systems-biology software 104 to verify, model, or analyze, for example, relative sequences, 3-dimensional structure, molecular



interactions, or overall cellular and physiological environment.

Systems-biology software 104 processes information and  
5 determines treatment dynamically for individual real-time  
physiological condition. Analysis report and other patient  
instructions are transmitted remotely as telemedicine  
service to network 103, which provides tasks to components,  
such as pharmaceutical or biopharmaceutical reservoirs 109,  
10 reconfigurable biosensors 102, wireless telemetry system  
106, therapeutic manufacturers 108, or other applications.

Sensors 102 may be hardware-reconfigurable or  
software-programmable according to user or systems-biology  
15 programming or report instructions. Ongoing or intermittent  
scheduled or random sensing events occurs between  
therapeutic components and pre-programmed and reconfigured  
micro/nano biosensors 102, along with proactive or reactive  
feedback to patient or user from systems-biology platform  
20 104. Preferably sensing process employs micro or nanoscale  
sensor 102 structure for minimal intrusion to individual  
health or physiology.

Optionally sensor system 100 provide wireless (RF)  
25 signal coupling with other sensors 100, such that  
communication occurs between different organisms having  
sensor 100. For example, sensor 100 may be implanted in  
pregnant host and another sensor 100 implanted in such host  
baby. Communication between sensors 100 may provide  
30 effective biological sensor signal transmission between  
separate hosts or organisms. Sensor 100 may be accessible  
according to IEEE 1451 network interface format.

Another example for multi-host communication implements sensors 100 for communication between separate related individuals, such as potential sexual partners, where one partner sensor 100 may sense sexually transmitted disease (STD) in such host, then such information is provided electronically to other host sensor 100 to produce proper antigens and antibodies to combat the STD.

Sensor unit 111 uses positioning device or chip 101 to position, locate or immobilize effectively target sample for analysis or sensing. The manipulated target sample comprises a biological molecule, organic or inorganic substance, such as cells, tissue, nutrients, chemicals, intracellular materials, extra-cellular materials, charged ions, pharmaceuticals, or molecular materials affecting host physiology.

Sensor unit 111 comprises multifunctional biosensor platform 102 for sensing and monitoring multiple biological materials, concentrations, inorganic or organic materials, cellular material, genetic material, nucleic acids, proteins, amino acids, peptides, antibodies, antigens, fatty acids, lipids, steroids, neurotransmitters, inorganic ions, pH levels, free radicals, carbohydrates, chemicals, small molecules, cells, tissue, pharmaceuticals, toxins, metabolites, or physiological levels macroscopically, microscopically, or nanoscopically.

Controller 112 uses network 103 to couple components for signal or data communication. Network 103 communicates data electronically to systems-biology platform 104.

Controller 112 may be implemented using personal, desktop, server, notebook, mainframe, wireless portable or other computer or processing device having processor, digital memory and network or user interface.

5

Systems-biology platform 104 uses computer equipment, software programs or reconfigurable firmware or emulation logic devices to verify, model, simulate, or analyze stored or raw data using computational biology, such as  
10 bioinformatics, proteomics, metabolomics, pharmacogenomics or other analysis software or hardware tools. Systems-biology platform 104 interprets or integrates data from biosensor platform 102, and analyzes organism preferably as a whole on system level. Systems-biology platform 104 may  
15 be integrated within one or more integrated circuit, module or processor; or bilaterally communicate to outside non-host signal source through wireless communication unit 106.

Controller 112 may use data storage 105 for storing  
20 processed data or applications programs from systems-biology platform 104. Controller 112 includes wireless communication unit 106, allowing bilateral communication with outside source, which may access or control sensor unit 111, controller 112, or therapeutic unit 113 through  
25 wireless communication unit 106.

Network 103 may couple therapeutic unit 113 with controller unit 112. Therapeutic unit 113 includes therapeutic manufacture 108 for providing pharmaceuticals,  
30 biopharmaceuticals, bio-catalytic chips or devices, tissue, or physiological treatments. Biopharmaceuticals include biological material for therapeutic use.

Therapeutic unit 113 includes therapeutic reservoir 109, which provides micro or nano-scale reservoirs containing pharmaceuticals or biopharmaceuticals. Contents  
5 of therapeutic reservoirs 109 may be provided or configured before sensor 100 is implanted in or attached to organism, or may be manufactured and filled *in vivo* by therapeutic manufacture 108. Therapeutic reservoirs 109 may release or dispense contents when appropriately signalled by network  
10 103.

Therapeutic unit 113 includes sensor manufacture 110 unit, which may provide additional sensors *in vivo* for additional targeted sensing or monitoring. Sensors from  
15 sensor manufacture 110 are part of or comprise biosensor platform 102.

Figure 1b shows positioning chip 101 for immobilizing or positioning target or tissue samples on or in sensor 102  
20 for bio-sensing as described herein. Positioning chip 101 may use micro-fabrication, micro-fluidics, or microbiology to manipulate, sort, or prepare samples, reagents, or other biological entities for analysis, high-throughput assays, or diagnostic applications. Positioning chip 101 may  
25 accomplish sample placement using multi-channel patch clamp electrophysiology chip to control individual cells by applying current to cell ion channels, positioning cells onto planar patch clamp, for example, e.g. Aviva Bioscience technique. The cell is sealed on-chip and analyzed or  
30 broken, and intracellular materials extracted and analyzed; if the cell is not analyzed, cellular material may be

positioned for analysis by diffusion, other natural technique, or through micro-fluidic manipulation.

Optionally, positioning chip 101 comprises microelectronic array or microfluidic assay, including  
5 electrodes or biosensors in which at least one microelectrode or sensor cavity or element is capable of generating controllable electric current or voltage for drawing probes, samples, or reagents to locations on sensor platform 102, allowing faster, controlled hybridization or  
10 analysis.

Positioning chip 101 may use micro or nano-chips with nanoscale channels or membranes, e.g., iMEDD NanoPORE membranes. Depending on size of such membranes, pores  
15 selectively exclude antibodies or proteins, while allowing free exchange of glucose, nutrients, insulin, or other molecules. Positioning chip 101 may position mammalian cells of host organism, as well as bacterial, fungal, protozoan, or other unicellular or multi-cellular organisms  
20 for analysis.

Additionally positioning chip 101 may detect or collect micro-metastatic tumor cells circulating in the blood stream or other body fluids, including but not  
25 limited to nipple aspirate, cerebrospinal fluid, peritoneal wash, sputum or excrement such as urine and stool. Preferably enrichment of tumor cells from blood stream may occur in miniaturized or microelectromechanical (MEMs) version of device such as autoMACS to collect circulating  
30 carcinoma cells from blood of patients with urologic cancers, or similarly using nanoparticles conjugated with

antibody to Epithelial Cell Adhesion Molecule to enrich for circulating tumor cells (CTC) of epithelial origin.

Further using positioning chip 101 in detection or  
5 collection, circulating prostate cancer cells in peripheral  
blood may be enriched, e.g., using technique by OncoQuick  
in Greiner, Germany, by using anti-human epithelial antigen  
paramagnetic microbeads or enrichment for disseminated  
breast cancer cells using advanced density gradient  
10 centrifugation; circulating endothelial cells serve as  
marker for vessel formation and vascular damage in cancer  
patients, such circulating cells being detectable for  
collection from peripheral blood using immunomagnetic beads  
coupled to antiCD146, an antibody raised against human  
15 umbilical vein endothelial cells.

Preferably collected tumor cells are analyzed on  
biosensor platform 102; for example, disseminated breast  
tumor cells may be analyzed by multiplex real-time RT-PCR  
20 (reverse transcriptase polymerase chain reaction) for  
mammoglobin, gabaII, B305D-C and B726P, or polymorphisms in  
carcinogen detoxifying UDP-glucuronosyl transferase UGT1A7  
in blood of patients with cancer of proximal digestive  
tract. Also enriched, using anti-epithelial cells antibody  
25 Ber-EP4, e.g., Dynal Corporation technique, epithelial  
cells derived from peripheral blood of prostate cancer  
patients can be analyzed using nested RT-PCR-PSA (reverse  
transcriptase polymerase chain reaction prostate specific  
antigen) assay as sensor mechanism.

30

Biosensor platform 102 may employ twenty-five  
epithelial tumor cells in bone marrow and lymph nodes of

esophageal carcinoma (pT1-3, pN0-1 and pM0) patients collected, using cytokeratin and EpCAM antibodies, respectively, by positioning chip 101 for micromanipulation in biosensor platform 102. Further DNA amplified by DNA  
5 sensor 201 using Mse-adapter PCR method may be analyzed by comparative genomic hybridization (CGH) for DNA-gains, -losses and point mutations by single-strand conformation polymorphism (SSCP). Also total RNA isolated PBMC in peripheral blood of breast cancer patients, may be subject  
10 to RT-PCR luminometric hybridization assay for presence of human telomerase reverse transcriptase, which is highly expressed in majority of tumor cells.

During sensing operation, positioning chip 101 may place  
15 samples on biosensor platform 102 for analysis. Biosensor platform 102 measures, detects, sequences, and other biological activities in serial or parallel in or out of organism. Biosensor platform 102 may use multi-functional high-throughput and density biochip having micro or  
20 nanoarrays, having substrates manufactured using glass, nylon, silicon, ceramic, metal, gel, membranes, synthesized nanomaterials, or other material.

Biosensor platform 102 provides data gathered from  
25 sensor arrays to network 103, which provides data to systems-biology platform 104, where data is integrated or processed. Systems-biology platform 104 may analyze empirically-sensed and simulated factors of individual organism in combination, to determine or confirm host profile of personal biological  
30 processes or makeup.

Systems-biology platform 104 may convey processed information to network 103. Network 103 communicates processed data to components coupled to network 103, including data storage 105, wireless communication unit 106, 5 therapeutic manufacture 108, therapeutic reservoirs 109, or sensor manufacture 110.

Data storage 105 keeps records or stores processed data by systems-biology platform 104. Processed data from systems- 10 biology platform 104, through network 103, optionally may be conveyed to wireless communication unit 106. Wireless communication unit 106 provides processed data access to external source, such as Global Positioning Satellite (GPS) receiver unit, media repository, personal computer (PC) or 15 workstation, laptop, handheld computing device, cellular device, internal or external camera, another internal implantable or attached sensor or chip, external biological monitoring device, outside network, healthcare provider, pharmacist, insurance agent, or other device or service 20 communicating with bio-sensor.

Processed data from systems-biology platform 104, through network 103, may be conveyed to therapeutic manufacture 108, where therapies are manufactured according 25 to host biological status or simulation output. Effectiveness or side-effects of therapies, produced by therapeutic manufacture 108, are monitored by biosensor platform 102. Ongoing or intermittent feedback from biosensor platform 102, through network 103, to therapeutic manufacture 108 provides 30 automated or iterative therapeutic process.



Optionally therapeutic manufacture 108 stores biological therapies in therapeutic reservoirs 109. Therapeutic manufacture 108 or therapeutic reservoirs 109 communicate through network 103 for filling or dispensing. Processed data  
5 from systems-biology platform 104, through network 103, may be conveyed to therapeutic reservoirs 109, where respective therapies are released according to biological status. Effectiveness or side effects of therapies, released by therapeutic reservoirs 109, are monitored by biosensor  
10 platform 102. For example, biosensor platform 102 may sense therapeutic effectiveness or side effects, while systems-biology platform 104 analyzes negative or positive effects to make recommendations. Ongoing feedback from biosensor platform 102, through network 103, to therapeutic reservoirs  
15 109 provides automated or iterative therapeutic cycle.

Processed data from systems-biology platform 104, through network 103, optionally is conveyed to sensor manufacture 110. Sensor manufacture 110 comprises hardware or  
20 software-programmable (reconfigurable and software-programmable terms may be used interchangeably) biosensors in vivo that integrate into biosensor platform 102 for supplementary sensing. Sensor manufacture 110 may be used to monitor additional biological materials originally part of  
25 biosensor platform 102, as well as used functionally to replace damaged sensors. Sensor manufacture 110 may be used to sense newly-calculated operational conditions by systems-biology platform 104. Optionally sensor manufacture 110 may monitor interactions between novel drug therapies, produced  
30 by therapeutic manufacture 108, and organism biology.

Appropriate timing of functions is preprogrammed before biosensor 100 is attached or implanted into organism. Time intervals for sensing are programmed according to external diagnosis, which can range from seconds, minutes, hours, 5 weeks, or longer. Once initial sensing begins, timing adjusts based on processed information by systems-biology platform 104. For example if genetic mutations within genome are found to be rare within multiplying cells, systems-biology platform 104 instructs biosensor platform 102 not to monitor genome as 10 frequently.

Conversely if sensed or simulation parameter, input vector, stimulus, condition, environment or other host biological factor is changing frequently, or there is a high 15 risk of change, then systems-biology platform 104 instructs biosensor platform 102 to increase frequency of particular sensor or assay. For example if organism changes through organ transplant, or is infected with new virus, systems-biology platform 104 instructs biosensor platform 102 to 20 increase the monitor frequency of antigen or antibody responses while decreasing such factors that are relatively stable.

Figure 2 shows biosensor platform 102 with 25 multifunctional array 200 coupled to detection system 230, and integrated sensor and detection system 231. Multifunctional array 200 serves as programmable or logical interconnect for coupling or switching various sensor devices, and interacts with samples and detection system 230 30 interprets samples into data to be analyzed by systems-biology platform 104. Multifunctional array 200 may include

micro and nanoarrays (M/N arrays) and biochips to test or monitor biological functions in particular organism.

Sensor components may include deoxyribonucleic acid (DNA) sensor 201, ribonucleic acid (RNA) sensor 202, peptide or protein sensor 203, antibody sensor 204, antigen sensor 205, tissue factor sensor 206, vector and virus vector sensor 207, lipid and fatty acid sensor 208, steroid sensor 209, neurotransmitter sensor 210, inorganic ion and electrochemical sensor 211, pH sensor 212, free radical sensor 213, carbohydrate sensor 214, neural sensor 215, chemical sensor 216, small molecule sensor 217, exon sensor 218, metabolite sensor 219, intermediates sensor 220, chromosome sensor 221, or cell sensor 222. M/N arrays are arranged architecturally as micro-electromechanical system (MEM) or as nano-electromechanical system (NEMS). This miniaturized architecture, as MEMS or NEMS device, allows multiple M/N arrays in a condensed form.

DNA sensor 201 is used to detect presence and/or sequence and/or structure of any DNA molecules including profiling for changes in methylation, monitor gene expression, undergo gene and DNA mapping, library screening, functional screen assays for nonsense and frame-shift mutations, scan the whole genome including micro-array-based comparative genomic hybridization to measure and map DNA copy number aberrations, detect disease markers, genotype single nucleotide polymorphisms (SNPs) including loss of heterozygosity analysis using SNP array hybridization and single-strand conformation polymorphism (SSCP), genotype organisms, examine protein-DNA interactions, and determine genetic characteristics individual to the organism.

DNA sensor 201 utilizes high-throughput M/N arrays for hybridization and use biochips, such as oligonucleotide M/N arrays, antibody M/N arrays, Pl-based artificial chromosome (PAC) M/N arrays, bacterial artificial chromosome (BAC) M/N arrays, yeast artificial chromosome (YAC) M/N arrays, cosmid M/N arrays, cDNA M/N arrays, gene M/N arrays, whole-genome M/N arrays, SNP M/N arrays, gridded cDNA M/N arrays, Southern Blots, theme M/N arrays (array centered around a particular disease or gene family), bead M/N arrays (arrays made up of small beads containing capture oligonucleotides), bead based M/N arrays (arrays in which reactions take place on the surface of microbeads), gel-pad M/N arrays (arrays in which chemical and enzymatic reactions can be carried out on three dimensional pads, like miniature test tubes), microcantilever arrays (in which specific biomolecular interactions occur on one surface of a cantilever beam, such as changes in intermolecular interactions that generate sufficient surface stress to bend beam for optical detection), M/N gel electrophoresis chips and M/N arrays 2D gel electrophoresis chips, chromatographic protein M/N arrays, e.g., Ciphergen protein sensor, and hybridization techniques for deoxyribonucleic acid sensing. Phenotypic markers for DNA damage or repair include single-cell gel electrophoresis use comet assay in which DNA damage is visualized, e.g., Komet 4.0 by (Kinetic Imaging Ltd) Imaging System.

Optionally for single nucleotide polymorphism (SNP) detection, DNA sensor 201 may apply so-called invader platform, or other device for genetic sequencing of an individual. DNA sensor 201 can analyze peritoneal fluid from

patients with ovarian cancer for loss of heterozygosity (LOH) at chromosomal arms 13 q(BRCA2 locus), 17 (BRCA1 and p53 loci) and 22q and for mutations in their p53 and k-ras genes. It can detect SNP (936 C>T) in 3' UTR of vascular endothelial growth factor gene (VEGF) in DNA extracted from blood of patients with breast cancer.

Further DNA sensor 201 can identify polymorphisms in carcinogen detoxifying UDP-glucuronosyl transferase UGT1A7 in blood of patients with cancer of the proximal digestive tract. Also methylation abnormalities in the promoter CpG islands of p16, HOX A9, MAGE A1 and MAGE B2 can be detected in sputum of lung cancer patients with DNA sensor 201. Sharply-elevated levels of stool DNA can be detected by DNA sensor 201 in patients with colorectal cancer. Stool DNA of surface epithelial cells is quantified using Picogreen fluorimetry.

DNA sensor 201 can detect chromosomal aneuploidy in cervical intraepithelial neoplasia or dysplasia using interphase cytogenetic technique called dual-color fluorescence in situ hybridization (FISH) targeting chromosomes 1, 7, 9 and 17 in Pap-smear slides and a thin layer of cervical cells.

Using DNA sensor 201, nipple aspirate fluid (NAF) containing epithelial cells shed from the breast ductal system can be analyzed. Extracted NAF DNA can be PCR amplified and analyzed for loss of heterozygosity in nuclear genome and deletions in mitochondrial genome using microsatellite markers and primer pairs, respectively.

Further DNA sensor 201 can be used to detect acute lymphoblastic leukaemia prenatally by analysing fetus blood to detect TEL-AML1 by FISH and genomic breakpoints by long-distance PCR. Using DNA sensor 201 and genomic DNA from whole  
5 blood, germ line polymorphism in KLK10 at codon 50 (GCC to TCC) associated with risk of occurrence in prostate cancer can be detected.

Also using DNA sensor 201, epigenetic changes, such as  
10 changes in GSTP1 methylation associated with prostate cancer can be detected in bodily fluids, e.g., urine and plasma, of prostate cancer patients. This detection uses real-time quantitative MSP and conventional MSP.

Further DNA sensor 201 is used to search for pieces of  
15 DNA in blood that are abnormally long, which is a signature of dying cancer cells; this test can be used for early diagnosis for patients with gynaecologic and breast cancers. Optionally oligonucleotide array-based genotyping platform,  
20 such as Perlegen, is used for accelerated SNP analysis, allowing whole-genome scanning by DNA sensor 201.

RNA sensor 202 may be used to detect presence, sequence or structure of RNA molecules, such as spliced and un-spliced  
25 RNA, mRNA, tRNA, rRNA, improperly transcribed RNA, properly transcribed RNA from diseased DNA sources, ribozymes, RNAi mechanism and application in relation to cancer therapy, or changes or differences in mRNA levels, or structures made of ribonucleic acids. RNA sensor 202 utilizes high-throughput  
30 M/N arrays for hybridization techniques, inclusive of DNA sensor 201. Probes may be made to hybridize with RNA

molecules, and Northern blot may be used in place of Southern blot technique.

RNA from enriched epithelial cells using anti-epithelial  
5 cells antibody Ber-EP4, e.g., per technique by Dynal  
Corporation, derived from peripheral blood of prostate cancer  
patients is analyzed for using nested RT-PCR-PSA assay by RNA  
sensor 202. Further, RNA sensor 202 can be used instead of  
second-look laparotomy in women with ovarian carcinoma  
10 treated with surgery and chemotherapy and show no sign of  
disease. Processed peritoneal washings are analyzed by  
telomerase repeat amplification protocol (TRAP) assay to  
detect residual disease. Total RNA isolated PBMC in  
peripheral blood of breast cancer patients, subjected to RT-  
15 PCR luminometric hybridization assay for presence of human  
telomerase reverse transcriptase that is highly expressed in  
majority of tumor cells.

Peptide or protein sensor 203 is used to detect primary,  
20 secondary, tertiary, or quaternary structures or activity of  
amino acid-based structures, such as sequence, enzymatic  
activity, protein function, interactions with agonists and  
antagonists, interactions with organic or inorganic  
structures or molecules, interactions with membranes, folding  
25 and enzymatic changes resulting in external factor, such as  
temperature, pH, ion concentrations, etc., N or C terminal  
characteristics, prions and misfolded proteins, amount and  
concentrations of proteins, bound and unbound state of  
proteins, sub-cellular localization, phosphorylated and  
30 dephosphorylated states, stages of degradation by proteases,  
stages of translation, gene and protein expression levels,  
e.g., using techniques such as ANTIBIOMIX (Milagen, Inc.) or

Antigen Retrieval (Biogenex Laboratories, Inc.), protein-protein interactions, protein-small molecule interactions, protein-antibody interactions, protein mutations due to transcription and translation mistakes, or measurable factors associated with amino acid based structures. Sensor 203 may be implemented using electrophoresis tag or microassay to identify protein or gene simultaneously, e.g., Aclara eTag assay (Mountain View, CA).

Peptide or protein sensor 203 utilizes high-throughput M/N arrays for hybridization and use biochips, such as protein M/N arrays, proteome M/N arrays, whole-proteome M/N arrays, electrospray fabricated protein M/N arrays, gene expression M/N arrays, reverse transfection M/N arrays (for example membrane proteins that are difficult to purify), functional protein M/N arrays, Western blotting, microcantilever arrays, or quantitative and qualitative high-throughput techniques for amino acid entities.

Peptide or protein sensor 203 can be used to detect proteins in cerebrospinal fluid of patients with primary brain tumors. Differentially-expressed proteins in processed CSF are digested and peptides identified by mass spectrometry. Presence of tumor-related proteins such as VEGF and VAV signifies presence of a primary brain tumor (179). Sensor 203, like SELDI protein-chip, similarly may be used to identify sixteen protein biomarkers in urine of bladder cancer patients, or instead of second look laprotomy in women with ovarian carcinoma who have been treated with surgery and chemotherapy and show no signs of disease. Processed peritoneal washings may be analyzed for telomerase activity to detect for residual disease.



Protein or peptide sensor 203 may be used in detection of diminished levels of N-CAM of <130 kDa in human serum of patients with brain tumors and the 80 kDa form associated with glioma. Further, protein and peptide sensor can be used in diagnosis of breast cancer by analysis of nipple aspirate fluid (NAF). Using SELDI-TOF capability, the presence of peptides at 4233.0 Da and 9470.0 Da is associated with cancer and the presence of 3415.6 Da and 4149.7 Da may be expected for normal samples. Thus sensor 203 can differentiate between diseased and unaffected populations.

Similarly protein sensor 203 may be used in breast-cancer diagnosis by analysis of serum samples. Samples applied to metal affinity capture chips activated with  $\text{Ni}^{2+}$ . Using SELDI protein chips/ mass spectrometry feature and software to detect selected discriminatory peaks separate cancer from non-cancer groups.

Using same features of sensor 203, serum is analyzed to differentiate between hepatocellular carcinoma (HCC) and chronic liver disease (CLD), where  $\alpha$ -fetoprotein fails as biomarker. Detecting 151 potential biomarkers in this way, system can provide diagnosis method for HCC. Using protein sensor 203 in diagnosis of prostate cancer, protein of 50.8kDa can be detected in serum even where PSA levels are <4ng/mL.

Further protein sensor 203 may be used in diagnosis of colorectal cancer detecting elevated HER-2 levels using standard ELISA and immuno-histo-chemistry (IHC) techniques. Elevated levels of secreted urokinase-type plasminogen

activator (uPA) can be detected by sensor 203 in serum for diagnosis of pancreatic cancer using sandwich ELISA or similarly, elevated levels of kallikrein 10 in serum for diagnosis of ovarian cancer, or elevated levels of basic fibroblast growth factor (bFGF) in nipple aspirate fluid in diagnosis of breast cancer, or elevated levels of fibroblast growth factor-2 and pleiotropin in serum for testicular cancer diagnosis or interleukin 6 in the serum of hormone-refractory breast cancer patients using immunoassay.

Antibody sensor 204 may be used to detect monoclonal or polyclonal antibodies. Similar to above sensors, hybridization with M/N arrays may be used. Probes may be chemical or molecular biological material that hybridize to targeted antibody, such as DNA, RNA, peptide, protein, small molecule, steroid, or lipid. Microcantilever arrays and other binding techniques can be applied.

Antibody sensor 204 may use so-called phagotope biochip to display phage with epitopes that react with antibodies in sera of patients with ovarian cancer, or other cancers. Also presence of elevated levels of anti-survivine autoantibody in serum of head or neck cancer patients is detected by antibody sensor 204 using recombinant protein survivine as antigen.

Antigen sensor 205 may be used to detect or recognize individual immune response factors. For example antigen sensing may detect autoimmune response factors, such as sensing multiplex character autoantibody response in systemic lupus erythematosus, rheumatoid arthritis, or multiple sclerosis. Another example of antigen sensor 205 application

may be identification or targeting of cell surface antigens for cancer therapy, e.g., Genentech approach.

Antigen sensor 205 may be used for early diagnosis of lung cancer or efficacy of chemotherapy by detecting nucleosomes in serum using assay, e.g., Cell Death Detection ELISApplus (Roche Diagnostics). Further antigen sensor 205 may detect tumor-associated antigens such as CYFRA21-1 for non-small cell lung cancer, and CEA, NSE and ProGRP for small-cell lung cancer.

Other sensing techniques for cancer detection contemplated herein include anti-malignin antibody screen test and tests for cancer markers including alpha fetoprotein (AFP), CA 15.3, CA 19.9, CA125, carcinoembryonic antigen (CEA), EVP test for Epstein bar virus, T/Tn Antigen test, TK-1 test and prostate specific antigen (PSA) or free PSA (fPSA) test. For bladder-cancer bladder-tumor-associated antigen test (BTA), BTA stat test, BTA TRAK test, fibrin/fibrinogen degradation products test (FDP), and NMP22 assay. Protein-based markers may illuminate and map abnormal cells, e.g., Inpath system. Other blood tests include CBC blood test, biological terrain assessment (BTA), Pre-Gen 26, telomerase test or DR-70 test.

Tissue-factor sensor 206 may use tissue factor M/N array to sense tissues, tissue factors, or tissue origin, using probes or antibodies to hybridize with targets. Tissue-factor sensor 206 may detect increase in prostaglandin E<sub>2</sub> production in cells that over-express COX2. This detection is associated with enhanced growth, migration and invasion as in bladder tumors.

Lipid or fatty acid sensor 208 may provide membrane mapping, M/N gel electrophoresis chips and M/N arrays 2D gel electrophoresis chips, detergent analysis, M/N array analysis  
5 of glycolipids and membrane proteins, membrane fluidity analysis, cholesterol analysis, or other test to examine cellular or intracellular organelles lipid bilayers.

Lipid or fatty acid sensor 208 may detect changes in  
10 exposed membrane; for example, such sensor 208 may produce antibody, with traceable label conjugated thereto, to anionic phospholipids (AP), such as phosphatidylserine, phosphatidylinositol and phosphatidic acid, that are more specific for AP than annexin V. When released into blood  
15 stream this antibody binds activated, by inflammatory cytokines, hypoxia, hydrogen peroxide, thrombin or acidic conditions, endothelial cells and thus, tumor blood vessels have increased exposure of anionic phospholipids on their surface. Localization of label enables localization of tumor.  
20 .

Lipid or fatty acid sensor 208 may detect levels of accumulation of synthetic membrane-permeable alkyl-lysophospholipids (ALPs), such as Edelfosine, Mitelfosine and Perifosine, that are anticancer agents that interfere with  
25 lipid mediated signal transduction.

Vector or virus vector sensor 207 may use microarray or assay with known sequenced virus attached e.g., DeRisi Laboratory. Unknown viruses may be detected through examining  
30 homology to known viruses, and subsequent arrays can be manufactured by sensor manufacture 110 to detect new viruses. Optionally assays that detect homologs can be applied, such

as Celera Diagnostic Viroseq™ HIV system for detection of mutations in human immunodeficiency virus (HIV) genome that confer drug resistance. Optionally assays for virus RNA can be used, such as Bayer Diagnostic Versant® HIV-I RNA 3.0 Assay for qualification of HIV-I RNA in plasma of infected people.

Further microparticle enzyme immunoassay (AxSYM HbsAg V2), e.g., Abbott Laboratories, may be used in quantifying reactivation of HBV during chemotherapy for lymphoma with Doxo rubicin along with real-time quantitative PCR specific to region of major S protein. Virus and virus vector sensor 207 may be used for detection of oncolytic virus replication in tumor tissues.

15

Steroid sensor 209 detects levels of steroids in the body, and monitors or controls levels of steroid hormones. Sensor 209 targets hormonal changes associated with puberty, menopause as well as fitness-conscious steroid-pumping athletic types.

20

Neurotransmitter sensor 210, small molecular sensor 217, or exons sensor 218 detects using M/N arrays, such specific antibodies as probes that hybridize with desired targets. Inorganic ion or electrochemical sensor 211 may detect ionic concentrations using techniques, using MEMS technologies with dielectric currents, microfluidics, or dialysis on a N/M platform. pH sensor 212 may read pH by detecting  $\text{H}_3\text{O}^+$  concentrations like silicon oxide pH sensors, e.g., Intelligent Pill. Free radical sensor 213 may be used to measure free radical activity, by using antioxidants as probes.

25

30

Carbohydrate sensor 214 may use oligosaccharide arrays, polysaccharide arrays, or carbohydrate chips, e.g., Glycominds glycochip, to measure glycan-protein interactions  
5 such as enzymes, antibodies, and lectins. Branched carbohydrates may bind to lectins involved in cell adhesion and migration processes. Also, natural branched carbohydrate like Lewis y, which is over-expressed in, for example, colon and ovarian cancer may be detected by carbohydrate sensor  
10 214. Such sensor 214 may apply to whole blood glucose (WBG) monitoring system, or continuous glucose monitor, e.g., Sensors for Medicine Science.

Neural sensor 215 measures action potentials or voltage  
15 between neurons in central nervous systems, using thin-film M/N electrodes as front-end sensors in MEMS and NEMS.

Chemical sensor 216 senses native or foreign chemicals, such as toxins, pharmaceuticals, vitamins, minerals, or other  
20 organic or inorganic chemicals. Chemical M/N arrays may be used, in which arrays of small organic compounds may be used to analyze interactions of proteins with various compounds. Conversely proteins or RNA may be used as probes to detect chemical substances.

25

Chemical sensor 216 may measure levels of carcinogen, benzo(a)pyrene diol epoxide, a metabolic product of benzo(a)pyrene found in tobacco smoke, known to cause 9p21  
abberations in peripheral blood lymphocytes in bladder cancer  
30 cases. Further chemical sensor 216 may measure tobacco-specific carcinogen 4-(methylnitrosamino)-1-(3-pyridyl)-1-

butanone (NNK) that can induce transformation of human breast epithelial cells, and may be directly related to initiation of human breast cancer in smokers.

Metabolites sensor 219 uses protein or antibody M/N  
5 arrays that hybridize to particular metabolites. Sensor 219 is useful to detect excessive buildup of metabolites. For example metabolites sensor 219 can measure serum homocysteine levels, associated with increased risk of cervical cancer, and further DNA sensor 201 may detect common polymorphisms in  
10 one-carbon metabolic pathway; examples of such mutations include MTHFR C677T, MTHFR A1298C and MTR A2756G. Increasing copies of MTHFR 677 variant polymorphism is associated with increased homocysteine levels whereas increasing copies of MTR 2756 variant polymorphism is associated with decreased  
15 levels of such metabolite.

Intermediates sensor 220 uses various protein and antibody M/N arrays that hybridize to particular intermediates. Sensor 220 is useful to detect excessive build  
20 up of intermediates; also sensing specific sequence, tertiary or quaternary structure of intermediates is used in drug design specificity.

Chromosome sensor 221 senses abnormalities in folding of  
25 chromosome, such as faulty histones, senescence-associated heterochromatic foci, or SAHF, since genes contained in these chromosomal regions are switched-on in proliferating cells, but are switched-off or "silenced" during cellular senescence.

30

Cell sensor 222 attaches whole living cells as probes, and is used for interactions with whole cells, such as cytotoxicity, drug metabolism, pharmacokinetics, target validation, interactions with other cells, extracellular materials, phenotypic analysis of genes and interfering RNA, as well as other biomolecules and compounds, e.g., Excellin Life Science bionic chip, which provides cell growth on chip. Effectively the cell becomes part of the chip, which allows manipulation and analysis of cell using microelectronics; the chip sends electrical signals through an on-board living cell, which detects changes in cell-membrane structure. The bionic chip can monitor and detect conditions that can cause cellular damage.

Optionally image cytometric measurement of breast fine needle aspirates can be used in cell sensing to predict nodal involvement in breast cancer DNA ploidy, S-phase fraction, G0G1/G2M ratio, and minimum (start) and maximum (end) nuclear pleomorphism indices (NPI). Further cytometric imaging allows differentiation between normal cells in which PML protein resides in discrete PML bodies and promyelocytic leukemic cells in which PML protein is genetically rearranged or dispersed throughout the nucleus.

Sensor unit 111 may measure or transmit blood pressure, flow rate or other sensor data wirelessly to controller unit 112, similarly to so-called cardioMEMS devices for monitoring pressure within aortic aneurysm. Biosensor 100 is implanted using catheter and transmits data to controller unit 112. Optionally such device can be used assessing circulation to organ after transplant or reconstructive surgery. This



provides physician with early indication of whether surgery is successful and prevent irreversible damage to organ.

5 Biosensor 100 may use implantable blood-flow monitoring system for providing synchronized blood vessel flow or myocardial wall contractility data to external monitor independent of transcutaneous leads. Further, since heart failure (HF) status of a patient is determined based on morphology of signal representative of arterial pulse  
10 pressure, the signal can be plethysmography signal that is produced by implantable or non-implanted sensor.

Time-derivative sensed signal may be produced based on signal representative of arterial pulse pressure; time  
15 derivation signal can be used to determine maximum and minimum peaks of signal representative of arterial pulse pressure. HF status can be accessed directly from time-derivative signal.

20 Biosensor 100 can be implanted using placement catheter, endoscope, or laparoscope; such device can be secured in LV or heart wall, e.g., using corkscrew, helical anchor, harpoon, threaded member, hook, barb, fastener, suture, or mesh or coating for receiving fibrous tissue growth.

25 Biosensor 100 provides less-invasive chronic measurement of left ventricular blood pressure or other parameters. Biosensor 100 can perform cardiosaver function to indicate to human subject that myocardial infarction is occurring; data  
30 is transmitted wirelessly to controller unit 112 for systems-biology analysis. Therapeutic reservoir 109 can inject thrombolytic or anti-thrombogenic agent into bloodstream

promptly to dissolve thrombus that caused myocardial infarction, and prevent formation of additional thrombi.

5 Biosensor 100 may sense impedance measurements of heart, respiratory or patient motion, and from these measurements, generating alarm signal when measurements indicate occurrence of cardiac arrhythmia. Optionally rate-responsive pacing system includes sensor of minimum oxygen content in right atrium over prescribed time interval, and using such minimum  
10 oxygen content as control parameter for adjusting rate of pacemaker.

Optionally for sensors in multi-functional array 200, nano-particles that specifically binds to particular  
15 molecules can be used to detect sequence, folding, binding interactions, function, or overall characteristics. Once bound to particular biological molecules, arrangement of distances between nanoparticles results in different observable properties, such as color or pattern.

20 Array 200 may be configured electronically by systems biology platform 104 to couple or interconnect selectively according to simulation or modelling to access actual host condition via one or more biosensor signals. Such sensed  
25 signal set may be compared by simulator against model or other software prediction to confirm host or target material health or other problem, as described herein.

Nanoparticle arrangement on biological molecules provide  
30 or indicate function, e.g., Northwestern University DNA-Driven Assembly of Biomaterials system. By attaching gold particles to DNA nucleotides, DNA hybridizes with

complementary strand and creates specific arrangement of gold particles. That arrangement of nanoparticles gives detectable color or pattern, which can be detected by optical device, and DNA can be sequenced.

5

Measuring color differences between nano-particle arrangement can also be applied to other biological molecule, e.g., Northwestern University Nanoscale Bioassay for Specific Antibodies. Rather than engineering nanoparticles that attach  
10 directly to the biological molecule, nanoparticles can be attached to specific antibodies. Binding of antibodies to targeted protein, DNA sequence, small particle, lipid, chemical, or other biological produces a particular color that is detectable or analyzable.

15

Also Nanoplex Technologies Nanobarcode Particles, made of different metals attached to biological molecules for multiplexing bioassays use probes attached to alternating metals on Nanobarcode to hybridize with biological molecules;  
20 then current can be run through Nanobarcode to determine molecules that bind to probes.

Detection system 230 may produce data from hybridization M/N arrays and other analysis techniques e.g., fluorescent  
25 scanners, laser scanning phosphorimagers, mass spectrometry, fiber optics, atomic force microscopy, parallel surface plasmon resonance imaging (allows direct analysis of binding events without need of reporter systems or tags), conclusive-induced dissociation (CID) mass spectra through electrospray  
30 ionization tandem mass spectrometry (ESI-MS) on triple or quadruple or ion trap mass spectrometers, real-time

polymerase chain reaction (PCR), PCR, Fluorescence *in situ* Hybridization (FISH), or charged coupled devices (CCDs).

Integrated sensor or detection system 231 may produce  
5 data from samples, without separating detection from  
hybridization or other technique. Optionally semiconductor-  
based M/N array can be used, e.g., CombiMatrix matrixarray;  
such array allows precise, digital control of electrochemical  
detritylation, including embedded sensor designed in  
10 semiconductor substrate, alternatively to conventional  
fluorescence technology. Hybridization with array sends  
direct electronic signals for analysis.

Another example of integrated sensor detection system  
15 231 assay can be GeneFluidics 3D micro-fabricated platform  
with embedded electrochemical sensor array. This platform  
conducts molecular analysis of raw DNA or protein samples,  
e.g., no PCR or immunoassays. Electrochemical detection of  
samples, such as whole blood, saliva, stomach acids, or other  
20 bodily fluids, uses current to measure electron transfer with  
current signal, associated with hybridized nanomolecules,  
e.g., ssDNA, hybridisable nanoparticles.

Biosensor 100 generally comprises biological  
25 microelectromechanical (bio-MEMs) sensor chip or detection or  
transducer device that may be implemented or computer-modeled  
for operation in silicon, silica, glass, polymer or other  
substrate or instrumentation cavity, beam, surface, channel,  
encapsulated molecules, membrane, quantum dot or nanocrystal  
30 (e.g., CdS, CdSe, CdTe, ZnSe, or other colloidal group II-VI  
semiconductor), matrix or array for single or multi-channel

independent signal detection in two or three dimensions in vitro or in vivo.

For example, sensor 100 may serve as high-throughput and  
5 sensitivity bio-physical, pharmaceutical or chemical  
recognition probe or cartridge for identification and/or  
characterization of host tissue or serum DNA, RNA, nucleic  
acids, protein, lipids, carbohydrates, enzymes, aptamers or  
other biomolecular or signal reporter target or any  
10 interaction, mutation, mass or rate thereof. Also such sensor  
may provide integrated, monolithic, discrete, or distributed,  
reagent-based or reagentless, microfluidic lab-on-chip  
microbiology mass spectrometry, flow immunosensor (e.g., FAST  
monitor for food or water quality), microarray or microassay  
15 functions, such as growing virus, bacteria or other  
eukaryotic or prokaryotic cells in microcells, nucleotide  
hybridization, polymerase chain reaction, molecular  
imprinting, chemical synthesis, ligand fishing, phage  
selection and concentration, multicomplex formation,  
20 diffusion limited concentration, or challenging antibiotics  
for rapid target detection, antibody susceptibility  
determination, or affinity and kinetic analysis.

Biosensor 100 may be implemented in quartz crystal  
25 microbalance for detecting or monitoring physical or chemical  
associated mass change or dissipation rate. Also whole cell  
or host sensor detection method may sense radioisotope,  
fluorescence, colorimetric, electrochemical,  
chemiluminescence, or bioluminescence. Additionally molecular  
30 or lipid-layer membrane-based sensor may operate to report  
change in electrical ionic, e.g., Ion Channel Switch  
biosensor using alternating current or voltage.

Furthermore encapsulated molecules may employ probes encapsulated by biologically localized embedding (e.g., PEBBLE nanosensors for intracellular chemical sensing, which  
5 may be delivered via gene gun, picoinjection, liposomal delivery, or phagocytosis, use matrices of cross-linked polyacrylamide, cross-linked decyl methacrylate, and sol-gel silica) for  $H^+$ ,  $Ca^{2+}$ ,  $K^+$ ,  $Na^+$ ,  $Mg^{2+}$ ,  $Zn^{2+}$ ,  $Cl^-$ ,  $NO_2^-$ ,  $O_2$ , NO and glucose detection; optionally encapsulated outer shell may be  
10 modified as configurable platform to target selectively specific biological locations or antibodies, such as including or excluding species variously reactive to passing through or filtered by the polymer membrane.

15 Biosensor 100 may recognize protein for antigen-antibody recognition, particularly by localizing or mapping protein residue epitopes. For example sensor contact at epitope-paratope interface functions via crystallographic analysis of one or more poly- or monoclonal or antigen-antibody complex.  
20 Also sensor 100 may detect cross-reactive binding with antiprotein antibodies using synthetic peptides as antigenic binding probe for free peptides or peptides adsorbed to solid-phase, conjugated to carrier or attached to synthesis support.

25 Additionally sensor 100 may detect cross-reactive binding decrease to identify critical residues in peptides via systematic residue replacement, as well as other protein-protein interaction, for example, between protease-inhibitor,  
30 antibody-antigen, enzyme-inhibitor, hormone-receptor, or signal transduction or transcriptional complexes. Protein sensing analyte may include fatty acids, maltose, biotin,

Ca<sup>2+</sup>, Co<sup>2+</sup>, Zn<sup>2+</sup>, Cu<sup>2+</sup>, glucose, glutamine, or other organic serum or tissue material.

5 Biosensor 100 may immobilize or control orientation of biomolecular target binding or catalytic sites via adsorption, entrapment behind membrane or in polymer or sol gel, covalent coupling, surface-immobilized polymer, or other capture system. Sensor orientation control may be accomplished via covalent coupling with attached glycosides, 10 generation of specifically-located thiol groups, use of antibody-binding proteins, avidin/streptavidin capture system, or use of tags with engineered antibody fragments.

15 Additionally sensor spatial control of surface immobilization may use soft lithography for substrate or surface patterning to introduce surface function, deposition control by physical placement, light-directed immobilization and patterning, or electro-chemical deposition control, for example, using elastometric polymer poly dimethylsiloxane 20 PDMS.

Also molecular imprinting polymer sensor may employ affinity sensor where response is produced by accumulation of template on sensor surface, receptor sensor where response is 25 generated by change in polymer characteristic or induced by template interaction, or enzyme-mimicking sensor where response is generated according to change in environment induced by molecular imprinting polymer-mediated catalytic reaction.

30

Furthermore antibody-based sol-gel sensor may use competitive assay detection, where antibody is encapsulated

in gel, sol-gel sensor is immersed in sample containing analyte concentration and known fluorescently labeled analyte solution, excess analyte is washed from gel, and fluorescence emission from remaining bound analyte is measured optically; 5 displacement assay detection, where antibody is encapsulated in gel with pre-bound fluorescently-labeled analyte, and gel is removed from solution and fluorescence emission from undisplaced analyte is measured; and fluorescence quenching detection, where fluorescently labelled antibody is 10 encapsulated in gel, which is immersed in sample, and bound analyte quenches fluorescence from antibody tag.

Biosensor 100 may employ optical biosensor or transducer with various assay formats. Direct assay may not use label, 15 and analyte surface binding is measured directly. Sandwich assay secondary antibody binds to surface-bound analyte molecule after analyte binding to sensor surface. Competitive assay enables binding-site competition on sensor surface, and low sensor signal is obtained for high analyte concentration.

20

Optical transducer sensor may use input grating coupler (e.g., bidiffractive grating coupler), prism coupler, planar or nonplanar, polarimetric, ion-exchange or deposited-rib, channelized or non-channelized waveguide or interferometer 25 (e.g., Mach-Zehnder interferometer), as well as surface plasmon resonance sensor (e.g., BIACORE system) using prism coupler, resonant mirror with vibro-stirrer (e.g., Iasys), evanescent wave fiber optic biosensor for multi-analyte detection (e.g., RAPTOR antibody identification system), 30 displacement flow detector, or other optical or time-resolved or phase fluorescence transducer (e.g., to detect



fluorophore-labeled binding protein or fluorescence resonance energy transfer), or fiber optic elements.

Biosensor 100 may employ acoustic transducer or wave  
5 device, such as bulk or surface acoustic wave device, thickness-shear mode resonator, shear-horizontal surface acoustic wave, acoustic plate mode, or love wave sensor, for example, to detect and characterize sensitive biological binding events in real time without labelling, by measuring  
10 energy loss occurring at liquid-solid biomolecular interface.

Biosensor 100 may employ fast-flow injection or microtiterplate immunoassay using enzymatic amplification electrodes, for example, via bi-enzymatic substrate recycling  
15 for signal amplification using electrochemical or bioelectrocatalytic redoxlabel immunoassay. Bioelectrocatalytic sensor electrode material for detecting phenolic targets via alkaline phosphatase measurement, for example, may include glassy carbon, graphite, carbon paste or  
20 ink, or gold.

Preferably sensing devices or techniques are provided or performed in miniaturized implantable format. However some sensor devices or methods may require sample from implanted  
25 device to be transferred to instrument located outside the body. Data generated by such instrument is transmitted to systems-biology platform 104 for analysis or modeling.

Biosensor platform 102 sensors, detection systems, or  
30 components may apply to parasitic or symbiotic organisms, such as bacteria, fungi, protozoa, plants, or other unicellular or multi-cellular organisms provided in host

organism, for example DNA sensor 201 may sense DNA structure of fungus cell living within such organism, peptide or protein sensor 202 may read its protein structures, and other sensors may read other biological properties. This  
5 information along with data from host organism is interpreted with systems-biology platform 104, and solution to expunge fungi is calculated or implemented.

Figure 3a shows software components of systems-biology  
10 platform 104. Once biosensor platform 102 produces comprehensive data on system, it is sent to network 103 and processed or analyzed by systems-biology platform 104.

Systems-biology platform 104 analyzes overall or partial  
15 structure of system or host, combining data from sensor components as well as model data of biosensor platform 102. Systems-biology platform 104 uses software for analyzing genomics 301, proteomics 302, computational chemistry 303, pharmacogenomics 304, computational biology 305,  
20 computational biophysics 306, computational cell behaviour 307, pharmacokinetics 308, metabolomics 309, transcriptomics 310, bioinformatics 311, other computational behaviour of the biological system, or other "omics" studies.

25 Other software may be integrated to understand or implement biological system on a personalized level, e.g., specific gene sequence, individual protein interactions, personal localized mRNA levels, dynamics of particular system, methods of control, personal cytotoxicity, and  
30 methods to design and modify the system; comprehensive data set is generated to understand fully or partially subject organism.

Genomics 301 may map, sequence, analyze, or discover function of organism genome. Structural or functional genomics may be used in genomics 301. Proteomics 302 analyzes organism proteome, describing set of proteins expressed during lifetime of cell or group of cells. Proteomics 302 calculates structure determination, at lower level, to functional analysis, or cell modeling at higher level of modeling.

Computational chemistry 303 uses algorithmic tools to facilitate chemical analyses. Chemical analysis occurs at atomic or molecular level, examining how individual and groups of atoms, compounds, or other structures interact with living system; further it analyzes chemical relationships between biological structures.

Pharmacogenomics 304 calculates potential drug responses based on personalized genetic information. This information is useful for determining appropriate therapies or preventing adverse reactions.

Computational biology 305 uses algorithmic tools to facilitate biological analyses. Computational biophysics 306 uses algorithmic tools to facilitate biophysical or biokinetic analyses. Computational cell behaviour 307 uses algorithmic tools to facilitate complete analyses of intracellular or intercellular behaviour.

Pharmacokinetics 308 determines or predicts kinetic interactions between potential drugs and organism biological molecules, taking into account variable interaction factors,

such as sterics, charge, dipole forces, or other factors that determine molecular interactions.

Metabolomics 309 analyzes organism overall metabolic profile; such as metabolism rates, amounts of metabolite intermediates, metabolic efficiency, structure of metabolic proteins, interactions between metabolic proteins and therapies, phosphorylative rates, or other aspects of individual metabolism.

Transcriptomics 310 analyzes organism transcription profile, such as efficiency, transcription errors to mRNA, intron-exon-splicing, biological transcription machinery, or other attributes of organism transcription.

Bioinformatics 311 undergoes database-management activities, involving persistent sets of data that are maintained in consistent state over indefinite periods of time. Bioinformatics 311 provides information content or flow in biological systems and processes; it serves as bridge between observations (i.e., data) in diverse biologically-related disciplines and derivations of understanding (i.e. information) about how systems or processes function, or subsequently the application.

Figure 3b shows ability to transfer information between systems-biology platform 104 and data storage 105 through network 103. This allows comparative studies between previously programmed and stored data with real-time computation; comparative studies serve as check against errors made by biosensor platform 102 and provide insights into overall systems understanding.

Also data storage 105 stores information processed by systems-biology platform 104. Data storage 105 may be located internally or externally relative to the organism, which can  
5 be accessed through wireless communication unit 106.

Regulation software or overlay 320 couples to data storage 105. When systems-biology platform 104 communicates with data storage 105, regulation overlay 320 assures that  
10 therapies, instructions, or other communications complies with Food and Drug Administration (FDA), Patent and Trademark Office (PTO), or other government regulatory bodies.

Regulation overlay 320 can store information or  
15 instructions for private agreements or regulations, such as contract or licensing agreement between biosensor 100 and pharmaceutical company. Depending on severity of organism condition or systems-biology platform 104 suggested therapy, communication directly or indirectly with FDA may be possible  
20 in instances where "expanded access," "compassionate use," "well characterized biological products," and other FDA exceptions apply. FDA may respond favourably and allow use of unapproved therapy (suggested by systems-biology platform 104) if exceptions apply.

25

Systems-biology platform 104 may implement neural network to model biological system or serve as decision aid for medical applications, problems or diagnosis. For example such platform 104 may employ methods as pattern recognition,  
30 feature extraction, supervised learning, unsupervised learning, or learning algorithms. Supervised learning methods may include Fisher's Linear Discriminant, Gradient Descent

Procedures, Perceptron Algorithm, Relaxation Procedures, or Potential Functions for linearly separable sets, or Nonlinear Discriminant Functions, Hypernet, Minimum Squared Error Procedures (MSE), or Ho-Kashyap Procedure for nonlinearly  
5 separable sets.

For multiple category classification problems, supervised learning methods may include the Fisher Linear Discriminant, Kesler Construction, or Backpropagation.  
10 Unsupervised learning methods may include clustering, Kohonen networks, Kohonen Competitive Learning, Hebbian learning, Adaptive Resonance Theory (ART) or prototype distribution map (PDM). Clustering approaches may include Basic Isodata Procedure, similarity measure approach, or criterion  
15 functions.

Criterion functions approaches may further include sum of squared error criteria, minimum error criteria, or scattering criteria, and such criteria may be used in an  
20 iterative optimization procedure. Platform 104 may also employ clustering approaches such as hierarchical clustering or metrics.

To assist in medical decision-making, systems-biology  
25 platform 104 may implement artificial intelligence or decision techniques, particularly data-based techniques or knowledge-based techniques. Data-base techniques may include approaches such as database, decision theory, pattern recognition, or Bayesian analysis, while knowledge-based  
30 techniques may include mathematical modeling and simulation, symbolic reasoning, as well as databases.

Systems-biology platform 104 may employ database such as patient record structures (e.g., hierarchical databases, National Library of Medicine, MUMPS (Massachusetts General Hospital Utility Multi-Programming System), ARAMIS system, PROMIS (problem-oriented medical information system), or medical database management system (e.g. MEDUS/A)). Systems platform 104 may employ disease database (e.g. oncology, rheumatology), or decision-support system (e.g. HELP program).

Platform 104 may employ differential diagnosis database (e.g. RECONSIDER or DXplain), online database, radiological database (e.g. CHORUS (collaborative Hypertext of Radiology)), or Human Genome Project. Mathematical modeling and simulation may apply to modeling of organism or biological process. Biological process may be represented by mathematical equations and evaluated.

Simulation involves representation of organism or biological process on a computer. Mathematical formulation may apply to administration of drugs or analysis of drug toxicity or drug level in a biological system. Pattern-recognition techniques may include discriminant analysis, method of classification using Bayes' Rule, parameter estimation, supervised learning or unsupervised learning.

Unsupervised techniques may include Parzen windworks, k-nearest neighbour estimation or other learning clustering techniques. Decision theory techniques may employ Bayesian analysis or Markovian analysis. Symbolic reasoning techniques may employ knowledge-based expert systems including early expert systems, second-generation expert systems. Techniques

of expert systems may include knowledge representation, heuristic search, natural language understanding, and exact reasoning. Second-generation expert systems may employ casual models, reasoning with uncertainty, or hybrid systems.

5

Systems-biology platform 104 may implement fuzzy techniques, (e.g. fuzzy set theory, fuzzy logic, fuzzy variables, or membership functions) for use in neural networks and expert systems. In dealing with uncertainty in supervised learning networks, neural networks may further employ pre-processing of fuzzy input, propagation of results through the network, or interpretation of final results.

Propagation of results may employ max-min networks, learning algorithms for interval data, or analogue models. Unsupervised learning methods may employ fuzzy associative memories or fuzzy clustering. Fuzzy methods for use in clustering include relation criterion functions, object criterion functions, fuzzy isodata, convex decomposition, numerical transitive, generalized nearest neighbour rules, or HCM/FCM clustering algorithm.

Uncertain information in knowledge-based system may employ fuzzy techniques when dealing with uncertainty in relation to input data, knowledge base, inference engine (e.g., binary logic engines or fuzzy logic engines), evidential reasoning (e.g., possibility theory, probabilistic approaches, or Dempster-Shafer Belief Theory), compatibility indices, or approximate reasoning.

30

Alternatively systems-biology platform 104 may employ probabilistic systems or statistical analysis for analysis of



medical data. Probabilistic systems may include Bayesian approaches, parameter estimation, discriminant analysis, statistical pattern classification, unsupervised learning, or regression analysis.

5

Bayesian approaches may include Bayes' Rule, Bayes' Decision Theory, risk analysis, supervised Bayesian learning, or decision trees. Parameter estimation may include maximum likelihood estimation or Bayesian estimation. Unsupervised learning may include Parzen window approach, nearest-neighbor algorithm, mixture densities and maximum likelihood estimates, unsupervised or Bayesian learning.

For example systems-biology platform 104 receives raw data from sensor unit 111 and employs neural networks, artificial intelligence, fuzzy systems, or probabilistic systems to aid in medical decision making for therapy recommendations or diagnosis.

Optionally additional information or test data helpful for diagnosis or treatment may be gathered from electronic files or user input from an outside source via and stored in data storage 105. Additional information or test data may include: patient age, height, weight, symptoms, allergies, diet, previous or present medications, medical or family history of disease, sickness or infection, results of previous blood, urine or other bodily fluid analysis, or other nongenetic (e.g., environmental) or immunological factors relating to the patient.

30

Optionally systems-biology platform 104 sends therapy recommendations or diagnosis report to an outside source via

wireless communication 106 and store recommendations or report in data storage 105.

In clinical, managed-care, hospital, diagnostic,  
5 therapeutic, or biomedical application or embodiment,  
systems-biology platform 104, using one or more firmware,  
source or object code software, configurable logic chip or  
device, digital signal processor, systolic processing array,  
or other finite state machine, actually or effectively may  
10 compare set of bioinformatic values associated with sensor  
signal or simulation data, preferably associated with same or  
different temporal states, to determine or otherwise  
recognize one or more genomic mutation associated with or  
corresponding to target patient, animal, plant, or other  
15 biological host.

Furthermore systems-biology platform 104 may operate  
autonomously, in cooperation with other computer system  
nodes, clients, or processing elements, to collect, process  
20 and display various host or patient sensor or simulation  
data, preferably in combination.

For example patient information and other personal or  
medical record data may be received via questionnaire or  
25 otherwise retrieved, such as host identification, drug  
treatment, prescription, and dosage, single or multiple  
concomitant food or drug allergy, interaction or side effect,  
pregnancy, lactation, as well as bioinformatic, genetic,  
proteomic, metabolomic, and other monitored, simulated or  
30 sensed mutation-related data as described herein.

Systems-biology platform 104 may be used in time-critical emergency, urgent, or trauma situation to improve patient health-care diagnosis and treatment, for example, by early-detection, expediting and assisting physician, paramedical, nursing or other professional analysis and treatment.

Sensed signal or simulated data electronically may be labeled for indicating genomic mutation, significantly improves quality and accuracy of medication delivery and administration to identified subgroups of patients having certain adverse response to medication, food, or other treatment.

Additionally such data or signal may include pharmacogenomic or pharmacokinetic clinical or indications based on genetic, proteomic, metabolomic (i.e., analysis of small organic cell molecules and metabolic response thereof), or other bioinformatic variant or mutation, or other genetic-based condition or profile (e.g., sex, race/ethnicity, etc.) such as drugs to be avoided, or considered as alternative. Thus, optimally, host susceptibility or predisposition to toxicity or other adverse host reaction or side effects to certain identified food, drugs, or other medical treatment may be minimized, mitigated, or eliminated using automated rule-based advice or expert system.

For example, systems-biology platform 104 may alert medical professionals when host patient is determined via sense or simulation approach to detect genomic mutation that patient ability to produce thiopurine S-methyltransferase (TPMT) enzyme activity is compromised. TPMT genetic test

(commercially available from DNA Sciences (Raleigh, NC)) enables identification of patient at risk for 6-MP/azathioprine/thioguanine toxicity, and improves confidence through tailored dosing regimens, while minimizing concern  
5 over drug-induced complication.

Alternatively, genomic mutation to G protein-coupled receptors (GPCR), molecular target and variant alleles may be detected to electronically label and thereby effectively  
10 modify host drug therapy. Another genomic mutation that may be detected and labeled is enzyme debrisoquine hydroxylase (CYP2D6), isozyme of microsomal cytochrome P450 monooxygenase system; encoding gene is located in CYP2D gene cluster in contiguous 45-kb region of chromosome 22. Here, at least  
15 nine polymorphisms of CYP2D6 affect metabolism of more than 30 different pharmaceuticals, including  $\beta$ -adrenergic receptor antagonists, neuroleptics, and tricyclic antidepressants.

Systems-biology platform 104 may couple electronically  
20 or digitally to hospital, physician, nursing, or other medical staff communication system to enable network-accessible prescription renewal, appointment scheduling, lab-result entry or retrieval, referrals to specialists and disease management, as well as generally computerized  
25 physician or pharmacy-ordering scheme, patient communications, access to medical simulation, test or sensor results, insurance claim status, and bar-coding of pharmaceuticals, and automated medication checks for possible errors.

30

System-biology platform 104 may employ simple identical or substantial equivalent value check between recently-

measured value and previously-stored value for same host, for example, after host exposure to radiation or other carcinogenic sources.

5        Such algorithm may be executed to adapt iteratively or dynamically in real-time or in multiple or parallel processors based on currently or recently-measured, monitored, or sensed host bioinformatic values, for example using fuzzy system, Bayesian or neural network, to improve  
10       compute or processing performance by comparing initially values that previously are known or recorded to be related or likely to be related or otherwise weighted to sensor signal or simulation data.

15       Additionally electronic access to sensor signal or simulation data may be restricted, secured, encrypted, or excluded unless the host thereof explicitly or voluntarily provides prior informed consent to access such information.

20       Hence, comparison serves to detect presence or absence of target sensor signal, simulation data or other genomic or bioinformatic value (e.g., oncogene, tumor suppressor gene, allele, enzyme, repeat sequence, micro-deletion, or other mutant gene product, protein, or metabolome) that causes, or  
25       increases or decreases risk of one or more host disease, disorder, syndrome, allergy, or other biological condition.

      Such simulation data or sensor information may be stored in data storage 105 or in other digital storage accessible or  
30       otherwise retrievable through network 103. Such stored information may be formatted according to one or more conventional, industry-standard, or otherwise publicly or

commercially-available software, processing, storage, and communications protocol, as well as databases for metabolic, signaling, regulatory and pathway data.

5        Additionally other genomic relational or object-oriented knowledge base or data sources may be network-accessed, such as GenBank, Unigene, LocusLink, Homologene, Ensemble, GoldenPath, or NCICB Cancer Genome Anatomy Project (CGAP). Such information may be accessed using ontology-based  
10 interfaces that are defined to be logically related, for example, using annotation format such as Distributed Annotation System (DAS).

Optionally systems-biology platform 104 data or  
15 instructions may be specified and otherwise annotated, such as hypothesis definition, experiment design, sample preparation and distribution, experiment run, data acquisition, result analysis, data mining, design refinement, modeling, knowledge discovery, or project report.  
20 Additionally such functions may be applied to simulation data or sensor signal processed by software or hardware analysis tools, e.g., for pharmacogenomics, gene expression, high-throughput sequencing, or proteomics (functional or structural) use-case domains.

25

Preferably such stored information complies, at least in part, with data exchange and management framework and specifications provided by Interoperable Informatics Infrastructure Consortium (I3C), which technical and use-case  
30 model documents, and recommended implementations, as described on-line at <http://www.i3c.org/> are hereby incorporated by reference as appropriate herein.

For example, one or more I3C-compliant or recommended data format may be employed during operation of electronic label processor, as described herein. Accordingly simulation  
5 data or sensor signal may be accessed, and displayed or otherwise imaged using electronic display I/O hardware or software, for gel chromatography images, original data from biological arrays, arrays of time-series data from mass spectrometry, illustrative functional depiction of proteins,  
10 simple microscope images, patient records with medical images, derived data from multiple or time-series images, electrocardiograms, or original drawings and annotations to medical images made by examining professionals. On-screen search capability enables medical professional quickly to  
15 locate and interpret particular host simulation data or sensor signal, such as gene sequence, protein, enzyme, allele, or other related detail.

Network 103 access to various databases or other digital  
20 repository may couple in n-tiered architecture multiple client interfaces, serve components, back-end objects and data sources. For example, Linux-based, Netscape, or Microsoft Internet Explorer browsers or applications, e.g., based on Java, non-Java, Perl, C, C++, or other programming  
25 or development software, run on client nodes 60 may receive information, such as in various mark up language e.g., HTML, XML, etc., from back-end objects over conventional network messaging or transport protocol, e.g., hyper text transfer protocol (HTTP), TCP Internet Protocol, simple object access  
30 protocol (SOAP), file transfer protocol (FTP), IIOP, etc. additionally Universal Description Discovery Integration

(UDDI) registry and Resource Description Framework (RDF) agent advertising formats may be used.

Further genomic, proteomic, or metabolomic sequence  
5 analysis software tool, for example, (e.g., BLAST, TimeLogic)  
may be used by controller 112 to discover or characterize  
host genomic, proteomic, or metabolomic sequence, acquired  
and qualified from one or more sources, such as sensor unit  
111 or data storage 105. Thus, internal and external sequence  
10 and protein libraries may be updated and maintained, certain  
redundant, unqualified or external data being filtered for  
internal sequence processing. One or more target, putative or  
otherwise mutant gene or bioinformatic value is then  
determined and catalogued effectively by systems-biology  
15 platform 104.

Hypothetical function of such determined gene or value  
may be generated manually, automatically, or homologously by  
finding similarity to known or other prior values. Genetic,  
20 proteomic, or metabolomic analysis protocols and similarity  
analysis may be defined and selected, thereby enabling or  
constructing functional hypotheses to be generated,  
prioritized, or reviewed using sensor measurements or other  
host evidence.

25

Proteolysis sample preparation may be performed (e.g.,  
HPLC, gel electrophoresis), then mass spectroscopy or tandem  
MS analysis and compression, quantitization, and fragment  
size genome analysis for candidate prediction, proteome or  
30 metabolome comparison, and other quantitative analysis using  
modeling tools and databases.



Systems-biology platform 104 may receive data from sensor unit 111, and neural networks, artificial intelligence, fuzzy systems, or probabilistic systems consider presence of conditions in diagnosis of genetic disorders: point mutations, mutations within non-coding sequences, deletions and insertions, trinucleotide repeat mutations, autosomal mutations, gain of function mutations, loss of function mutations, mutations in mitochondrial genes, enzyme defects, defects in receptors and transports systems, defects in receptors and transport systems, alterations in structure, function or quantity of non-enzyme proteins, defects in receptor proteins, defects in protooncogenes or tumor-suppressor genes, aneuploidy, unbalanced autosome, sex chromosome abnormality, fragile X syndrome, ring chromosome, chromosome inversion, isochromosome formation, translocation, or abnormal gene products.

Optionally allele-specific oligonucleotide hybridization may be employed in multifunctional array 200 in biosensor platform 102 to assist in direct gene diagnosis of mutations. Systems-biology platform 104 may diagnose genetic disease or mutation, such as Mendelian disorders, autosomal dominant disorders, autosomal recessive disorders, X-linked disorders, Marfan syndrome, Ehlers-Danlos syndrome, familial hypercholesterolemia, lysosomal storage diseases, Tay-Sachs Disease, Gangliosidosis, Niemann-Pick disease, Gaucher Disease, glycogen storage diseases, Mucopolysaccharidoses, Alkaptonuria, Neurofibromatosis, trisomy 21, chromosome 22q11 deletion syndrome, Klinefelter syndrome, XYY syndrome, Turner Syndrome, Multi-X females, hermaphroditism, pseudohermaphroditism, triplet repeat mutations, chromosome-

breakage syndrome, Prader-Willi syndrome, Angelman syndrome, or gonadal mosaicism.

Alternatively, systems-biology platform 104 may diagnose  
5 infections disease or infection such as Haemophilus  
influenzae infection, tuberculosis, histoplasmosis,  
coccidioidomycosis, shigella bacillary dysentery,  
Campylobacter enteritis, Yersinia enteritis, Salmonellosis,  
typhoid fever, cholera, amebiasis, giardiasis, herpes,  
10 Chlamydia, gonorrhea, syphilis, trichomoniasis,  
staphylococcal infection, streptococcal infection,  
clostridial infection, measles, mumps, mononucleosis, polio,  
chickenpox, shingles, whooping cough, diphtheria, infections  
associated with Neutropenia and Helper-T cell depletion,  
15 cytomegalic inclusion disease, pseudomonas infection,  
legionnaires disease, listeriosis, candidiasis,  
cryptococcosis, aspergillosis, mucormycosis, pneumocystis,  
pneumonia, cryptosporidium and cyclospora infection,  
toxoplasmosis, dengue fever, Rickettsial Infection, trachoma,  
20 leprosy, plague, relapsing fever, lyme disease, malaria,  
babesiosis, leishmaniasis, African Trypanosomiasis, Chagas  
disease, Trichinellosis, hookworm, cysticercosis, Hydatid  
disease, chistosomiasis, lymphatic filariasis, or  
onchocerciasis.

25

In diagnosing infectious disease or infection, systems-  
biology platform 104 receives data from sensor unit 111 or  
neural networks, artificial intelligence, fuzzy systems, or  
probabilistic systems that consider presence of infectious  
30 agent, such as a prion, virus, bacteriophage, plasmid,  
transposon, chlamydiae, rickettsiae, mycoplasma, fungi,  
protozoa, helminths, or ectoparasite. In host system,

systems-biology platform 104 may also consider the presence of bacterial endotoxin, bacterial exotoxins, proliferation and morphologic lesions of epithelial cells, tissue necrosis, granulomas, cysts, increased levels of leukocytes, mononuclear cells or neutrophils, mononuclear interstitial infiltrates, reduced levels of immune cells (e.g. cytokines, lymphocytes, macrophages, dendritic cells or natural killer cells), bacterial leukotoxins, hemagglutinin, spores, or other antigen or protein from bacteria, virus, fungi, protozoa, or parasite.

Alternatively systems-biology platform 104 may diagnose disease of immunity, such as hypersensitivity disorders (immune complex mediated, complement-dependent reactions, cell mediated, or anaphylactic type, transplant rejection) autoimmune disease, systemic sclerosis, inflammatory myopathies, mixed connective tissue disease, polyarteritis nodosa or other vasculitides, X-linked agammaglobulinemia of Bruton, common variable immunodeficiency, isolated IgA deficiency, Hyper IgM syndrome, DiGeorge syndrome, severe combined immunodeficiency disease, immunodeficiency with thrombocytopenia and eczema, acquired immunodeficiency syndrome (AIDS), or amyloidosis.

In diagnosing immunity diseases, systems-biology platform 104 considers following sensed, detected, or measured conditions from sensor unit 111: levels of immune cells (e.g., mast cells, cytokines, lymphocytes, macrophages, dendritic cells or natural killer cells), MHC (major histocompatibility complex) molecules or antigens, HLA (human leukocyte antigen) complex, antigens, or types, or levels of primary mediators (e.g., biogenic amines, chemotactic

mediators, enzymes, or proteoglycans), secondary mediators (e.g., leukotrienes, prostaglandins, platelet-activating factors, or cytokines), histamines, platelet-activating factor (PAF), neutral proteases, chemotactic factors, or  
5 antigen-presenting cells (APC).

In diagnosing autoimmunity diseases, systems-biology platform 104 receives data from sensor unit 111 or neural networks, artificial intelligence, fuzzy systems, or  
10 probabilistic systems considers presence of auto-antibodies disease and considers whether auto-antibodies are directed against single organ or cell type or whether it is systemic. Autoimmune diseases include single organ or cell type related diseases (e.g., hashimoto thyroiditis, autoimmune hemolytic  
15 anemia, autoimmune atrophic gastritis of pernicious anemia, autoimmune encephalomyelitis, autoimmune orchitis, goodpasture syndrome, autoimmune thrombocytopenia, insulin-dependent diabetes mellitus, myasthenia gravis, Graves disease), or systemic autoimmune diseases (e.g., systemic  
20 lupus erythematosus, rheumatoid arthritis, Sjögren syndrome, or Reiter syndrome).

Systems-biology platform 104 may identify whether disease condition may be single organ or cell type autoimmune  
25 diseases or primary biliary cirrhosis, chronic active hepatitis, ulcerative colitis, or membranous glomerulonephritis. The platform is also identifies whether disease condition may be systemic autoimmune disease or inflammatory myopathies, systemic sclerosis (scleroderma) or  
30 polyarteritis nodosa.

Furthermore systems-biology platform 104 may determine presence of pathologic autoimmunity by considering at least three requirements, such as presence of autoimmune reaction, clinical or experimental evidence that such reaction is not  
5 secondary to tissue damage but of primary pathogenetic significance, or absence of another well-defined cause of disease.

Alternatively systems-biology platform may be used in  
10 diagnosis of neoplasia. In diagnosing neoplasia, systems-biology platform 104 receives sensed, detected, or measured data from sensor unit 111 and neural networks, artificial intelligence, fuzzy systems, or probabilistic systems considers the following factors: DNA damage, failure of DNA  
15 repair, mutations in the genome of somatic cells, activation of growth-promoting oncogenes, alterations in the genes that regulate apoptosis, inactivation of cancer suppressor genes, expression of altered gene products and loss of regulatory gene products, oncoproteins, growth factors, growth factor  
20 receptors, proteins involved in signal transduction, nuclear regulatory proteins, cell cycle regulators, tumor antigens, or the levels of immune cells (e.g., mast cells, cytokines, lymphocytes, macrophages, dendritic cells or natural killer cells).

25 Systems-biology platform 104 may consider epidemiological factors in determining diagnosis for neoplasia. Epidemiological factors may include cancer incidence, geographic or environmental factors (DNA damaging  
30 agents - e.g. chemicals, radiation or viruses), or heredity (e.g., inherited cancer syndromes, familial cancers, autosomal recessive syndromes of defective DNA repair).

Systems-biology platform 104 may consider tumor markers such as hormones (e.g. human chorionic gonadotropin, calcitonin, catecholamine and metabolites, or ectopic hormones), oncofetal antigens ( $\alpha$ -fetoprotein or carcinoembryonic antigen), isoenzymes (e.g., prostatic acid phosphatase, or neuron-specific enolase), immunoglobulins, prostate-specific antigens or mucins or other glycoproteins (e.g. CA-125, CA-19-9, or CA-15-3).

After systems-biology platform 104 makes diagnosis, platform may recommend treatments in combination or individually. Such recommendation may include diet changes, surgery, radiation therapy, chemotherapy, medications, antiangiogenesis therapy, or other cancer treatment. Systems-biology platform 104 may instruct therapeutic unit 113 to manufacture or dispense pharmaceuticals, biopharmaceuticals, or other therapeutic tools for the treatment of neoplasia.

Systems-biology platform 104 may employ sensor device and simulation method for analyzing dynamic hormone-secretion phenomena in dynamic biological systems, for example using sensor, artificial neural network, and dosing device; e.g., Sichel Technologies wireless or telemetric sensor platform for measuring parameters of relevance in vivo, such as radiation dose, tissue microenvironment or gene expression to increase treatment success. Implantable sensors may be provided 2mm diameter, 15mm length, for injection at margin of tumors using minimally invasive procedure.

Biosensor 100 may be applied to food technology, e.g., pasteurization or development or production of artisan foods. DNA sensor 201 may monitor, detect, or measure amount of

bacteria or microflora used to ripen and develop flavours in foods, such as artisan cheese. Similarly peptide or protein sensor 203, lipid or fatty acid sensor 208, or small molecule sensor 217 may monitor bacterial or microflora production of  
5 fats, proteins, esters, or other biologically-active molecules.

Biosensor 100 may be applied to food manufacturing industry, e.g., quality control, food safety, or countering  
10 food borne illness caused by bioterrorism. Biosensor 100 may detect types of food contaminants, including bacteria or chemicals that cause human sickness, or counter bioterrorism acts threatening consumer food supply.

15 Biosensor 100 may be used by food manufacturer, crop cultivator, lab researcher, consumer, packer, distributor, receiver, food vendor, or food inspector to ensure quality control and food safety. Biosensor platform 102 may detect, measure, or determine presence or absence of parasitic  
20 organism, virus, bacteria, fungi, protozoa, or unicellular or multi-cellular organism present during food manufacturing process or growth of food crops, or prior to consumption.

Chemical sensor 216 may be used to sense, detect, or  
25 measure foreign chemicals, such as toxins, vitamins, minerals or other organic and inorganic chemicals. Systems-biology platform 104 may analyze raw data from biosensor platform 102 to identify potentially-hazardous organism or chemical or flag unknown organism or chemical.

30

When systems-biology platform 104 identifies or quantifies potentially hazardous organism or chemical or

unknown organism or chemical, data is stored in storage 105. Systems-biology platform 104 may generate report document or electronic multi-media warning or signal, which discloses detected organism or chemical and determine whether  
5 manufacturing, crop growth, or consumption is safe to continue.

Systems-biology platform 104 may send automated warning or signal, sent via wireless communication 106, to  
10 information recipient interested in data gathered by the platform, such as remote database, researcher, lab, government agency, or health or safety maintenance organization.

15 Chemical sensor 216 may determine purity or verify amount of vitamin, mineral, herb, or botanical claimed by a food product, meal supplement, vitamin supplement, or other nutritional substance. Systems-biology platform 104 may compare amount of vitamin, mineral, herb or botanical  
20 determined by chemical sensor 216 to pre-set amount or range stored in storage 105, e.g. amount or range determined by a government agency or health or safety maintenance organization.

25 Systems-biology platform 104 generates report whether detected amount or range complies with pre-set amount or range, and determines whether manufacturing or consumption is safe to continue. Detected amount can be reported and sent via wireless communication 106 to outside source or  
30 information recipient interested in data gathered by chemical sensor 216, such as packer, distributor, receiver, remote database, researcher, lab, government agency, or health or



safety maintenance organization. During manufacturing, determined amount of vitamin, mineral, herb, or botanical present in each lot or batch of produced product is recorded or accessible through network 103 for analysis.

5

Optionally if amount of vitamin, mineral, herb, or botanical falls outside pre-set amount or range, systems-biology platform 104 generates automated warning to outside source or information recipient. Biosensor 100 monitors  
10 manufacturing of food product, meal supplement, vitamin supplement, or other nutritional substance by ensuring that manufactured substance complies with required amount or range of nutritional substance. Chemical sensor 216 may be used to demonstrate whether particular vitamin, mineral, herb,  
15 botanic, or other natural or organic food has properly absorbed in biological system of organism.

Biosensor 100 may synchronize different input stimuli, particularly with integrated purpose of evaluating food and  
20 drug interactions positively or negatively within host. Systems-biology platform 104 can analyze genetic composition of host, determined through DNA sensor 201, to assist in predicting particular drug-food interactions. To assist in predicting drug and food interactions, host genetic  
25 composition may be supplemented with additional information or test data including nongenetic (e.g. environmental, epidemiological) or immunological factors relating to host.

Biosensor 100 may be implanted within a host and  
30 pharmacogenetics 304 or pharmacokinetics 308 in systems-biology platform 104 may be employed to monitor or determine activity or effectiveness of medication used individually or

in combination. Meanwhile, biosensor 100 placed remotely or separately from implanted biosensor is used to analyze nutritional substance (e.g., food product, meal supplement, vitamin, or mineral) that may be consumed by same host.

5

Data from remote biosensor 100 is coupled, received, or combined to data from implanted biosensor or analyzed collectively by systems-biology platform 104 to predict or model combined allergic reactions, side effects, or adverse reactions that result from consumption of nutritional substance in conjunction temporally with related medication.

Systems-biology platform 104 may generate automated recommendation or report diagnostically or therapeutically about optimum level of nutritional substance or identify alternative substance for consumption. Data from remote and implantable biosensor data, and recommendation or determination processed by systems-biology platform 104 may be stored in data storage 105. An outside source or information recipient may access data and results in data storage 105 through wireless communication 106 for analysis via network 103.

When systems-biology 104 identifies nutritional substance that may cause an adverse or positive reaction, automated warning or message may be transmitted wirelessly to information receipt interested in the gathered data. The ability of systems-biology platform to analyze or model nutritional substance and host condition in combination using host sensor data and consumable sensor data optimizes treatment of real-time physiological condition.

Biosensor 100 may be applied to biopharming purpose, e.g., field tests or inspections of genetically engineered plants, and use of genetically engineered plants or transgenic crops to produce therapeutic proteins and industrial enzymes with safeguards for ensuring that food crops are not co-mingled with food crops intended for pharmaceutical or industrial use.

To prevent out-crossing or commingling of genetic material, DNA sensor 201, RNA sensor 202, or peptide and proteins sensor 203 in biosensor platform 102 may detect, sense or measure presence or absence of foreign genetic material or protein in food crop not intended for pharmaceutical or industrial use. Systems-biology platform 104 may analyze raw data from biosensor platform 102 to identify out-crossing or commingling of genetic material.

When systems-biology platform 104 identifies foreign genetic material, data is stored in storage 105. Systems-biology platform 104 may generate report about detected foreign genetic material or determine whether crop growth is safe to continue. Systems-biology platform 104 may send automated warning or signal, via wireless communication 106, to information recipient interested in data gathered by platform, such as remote database, researcher, lab, government agency, or health or safety maintenance organization.

Biosensor 100 may monitor growth of food crops e.g. sensors (e.g. peptide or protein sensor 203, vector or virus vector sensor 207, pH sensor 212, metabolites sensor 219, etc.) in biosensor platform sensor 201 may sense, detect or

measure abnormalities in crop growth or reproduction. Biosensor 100 may monitor, detect or measure pesticides, insecticides or foreign chemicals effect on growth or reproduction.

5

Biosensor 100 may be applied to bio-manufacturing industry, e.g., drug-producing plants and transgenic animals, such as cows genetically transformed to excrete different kinds of therapeutic proteins in breast milk. Peptide or protein sensor 203 in biosensor platform 102 or antibody sensor 204 may detect or measure presence or absence of genetically engineered therapeutic protein or antibody in breast milk or other biological fluid.

15 Biosensor 100 may be applied in xenotransplantation, for example by screening animal organs for transplantation into humans. Sensor unit 111 senses, measures, or processes biological molecule, such as cell, tissue, or intracellular or extracellular material from animal cell, tissue or organ, or raw data is analyzed by system biology platform 104. System biology platform 104 analyzes or determines whether animal cell, tissue, or organ is compatible for use with human for transplantation or other therapeutic process.

25 Biosensor 100 may be applied to avian transgenics, particularly to proteins produced through poultry-based production systems. For example, biosensor platform 10 may detect whether successful transformation is occurring via avian embryonic germ cell, retroviral-mediated transformation, sperm-mediated transgenesis, avian embryonic stem cell, direct egg transfection, or other transformation process.

Biosensor 100 may be applied to drug-producing plants, e.g., tobacco, corn, or other non-food plants, for biomanufacturing. Peptide or protein sensor 203 may detect, sense or measure presence, absence, manufacture or biological activity of recombinant proteins manufactured in plants. DNA sensor 201, RNA sensor 202, vector or virus vector sensor 207, chromosome sensor 221, or cell sensor 222 may monitor or detect whether genetic material, vector, chromosome, or cell successfully integrates or genetically transforms plant or animal.

Figure 3c systems-biology platform 104, therapeutic unit 113, and sensor unit 111. Systems-biology platform 104 provides verification of data 321, to assure that data is proper or feasible from biosensor platform 102 within sensor unit 111. Verification of data 321 identifies sequence or structures of target system. Data may be analyzed statistically by systems-biology platform 104, using statistical computation, e.g., scatter plot matrices, Venn diagrams, comparative histograms, volcano plots, or gene ontology charts. Computed statistics are interpreted biologically, filtering or reducing dataset to manageable size by eliminating results that show insignificant or uninteresting biological data.

Verification of data 321 includes checking regulatory relationship of genes or interaction of proteins that provide signal transduction or metabolism pathways, as well as physical structure of organisms, organelle, chromatin, cell-cell interactions, or other components.

To integrate sensor data, software and management systems are used. Systems-biology platform 104 may utilize management software, e.g., Analysis Information Management System (AIMS), using tools to analyze or manage range of complexity of data obtained from microarrays or assays, tracking computational processes. Data-mining tools, e.g., high-dimensional data analysis tools, may process data where data have multiple dimensions.

Data may be formatted using standardization programs, e.g., Gene Expression Markup Language (GEML), Microarray Markup Language (MAML), Microarray and Gene Expression Data (MAGE), MicroArray and Gene Expression Markup Language (MAGE-ML), solutions by Microarray Gene Expression Database group (MGED) or Minimum Information About a Microarray Experiment (MAIME), or other programs.

After data is verified, modeling/simulation 322 uses combined simulation data or sensor signal to model biological structures or relative interactions. Modeling or simulation 322 simulates biological interactions to identify behaviour of system, for example, sensitivity of behaviours against external perturbations and how quickly system returns to normal state after stimuli.

Another example includes simulating how individual malfunctioning mis-folded protein interacts with other proteins or cellular components, with simulations on how protein responds to particular therapies; yet another example is modeling phospho-proteomics and systems biological role for oncology target discovery or validation.

Modeling or simulation 322 predicts methods of  
controlling state of biological system, e.g., pharmaceutical  
or gene therapy transformation of malfunctioning cells into  
healthy cells. For example through structural analysis,  
5 regulation of c-Ab1 and STI-571 specificity may be achieved.

Modeling or simulation 322 prediction is translated into  
instructions for therapeutic unit 113 to implement  
appropriate therapy to fix biological system. These  
10 instructions are conveyed to therapeutic unit 113, where  
instructed therapy may be performed.

Sensing unit 111 monitors progress, efficiency, or  
ancillary effects of induced therapy on biological system.  
15 Data from sensing unit 111 may be verified by verification of  
data 321, which provides cyclical self-regulating process.

Figure 4a shows flow of instruction from systems-biology  
platform 104 to network 103 to components comprising  
20 therapeutic unit 113. Components of therapeutic unit 113  
include therapeutic manufacture 108, therapeutic reservoirs  
109, and reconfigurable sensor manufacture 110. These  
components may be reconfigurable or software-programmable  
according to systems-biology platform 104, or from external  
25 source through wireless communication unit 106.

Figure 4b shows therapeutic manufacture 108 of:  
pharmaceuticals 401, biopharmaceuticals 402, tissue,  
reconfigurable biocatalytic chips 403, tissue scaffolds 404,  
30 M/N machines 405, or other therapeutic material or tools.  
These components may be reconfigurable and software-

programmable according to systems-biology platform 104, or from external source through wireless communication unit 106.

5        Pharmaceuticals 401 may be known and matched with organism, or computationally derived or optimized from systems-biology platform 104. Pharmaceuticals 401 is defined herein as including chemical substance that provides benefit to system.

10        Biopharmaceuticals can be naturally-occurring biological molecule or structural derivative of biological molecule. For example, biopharmaceuticals can be isolated DNA molecules, recombinant DNA molecules, DNA fragments, oligonucleotides, antisense oligonucleotides, RNA molecules or constructs,  
15        self-modifying RNA molecules, catalytic RNAs, ribozymes, modified ribozymes, synthetic peptides, peptide linkers, proteins, fusion proteins, antibodies, modified antibodies, antigens, cell surface receptors, monoclonal antibodies specific for epitopes, polyclonal antibodies, tissue factors,  
20        modified tissue factors, mutant tissue factors, ligands, vectors, virus strains for gene transfer, recombinant plant viral nucleic acids, bacterial strains, oil-body proteins as carries of high-value peptide in plants, host cells, transformed cells, or microorganisms newly isolated in pure  
25        form from natural source.

      Therapeutic unit 108 may prepare biopharmaceutical product such as 2 vg of sub50-nm tenascin nanocapsules containing antisense of protein kinase CK2  $\alpha$  subunit or  
30        similarly GFP and RFP-labeled bacteria which produce toxins or other therapeutic proteins to be used to target tumors. Further therapeutic unit 108 can perform functions like so-



called Intelligent Pill (e.g., University of Calgary) in which information relayed to chip that controls micropumps that squeeze-out therapeutic material.

5           Therapeutic manufacture unit 108 may prepare therapy comprising pharmaceutical 401 or biopharmaceutical aspect 402. For example antiangiogenesis thereapy using yttrium-90 nanoparticles with conjugated anti-Flk-1 monoclonal antibody administered by i.v. injection is anti-angiogenic agent for  
10 treatment of solid tumors. Therapeutic manufacture unit 108 may produce small interfering RNA (siRNA) used to inhibit P-gp encoded by *MDR1* gene; production enhances accumulation of sensitivity of multidrug-resistant cancer cells to drugs transported by P-glycoprotein.

15           Reconfigurable biocatalytic chips 403 are software programmable from instructions by systems-biology platform 104, or from external source through wireless communication unit 106. Depending on instructions, reconfigurable  
20 biocatalytic chips 403 can be activated, deactivated, manufactured, or disassembled. Reconfigurable biocatalytic chips 403 undergo molecular bioprocessing, fabricating or manipulating single and multienzyme systems on biochip to induce artificially biocatalysis in system.

25           Tissue scaffolds 404 may be reconfigurable, and controlled by systems-biology platform 104 instructions (or from external source through wireless communication unit 106). Scaffold 404 may be substrate to grow cells or tissues,  
30 which may be activated or deactivated according to signalled instructions. Permanent or biodegradable tissue scaffolds can be used. Further scaffold 404 may be personalized by systems-

biology platform, e.g., John Hopkins University stem cell-based polymer scaffolds for tissue engineering using composite hydrogel. After modeling tissue development on biomaterial scaffolds based on individualized systems-biology profile, reconfigurable scaffold 404 can be programmed with biological signals based on individual need.

M/N tools 405 may perform therapeutic treatments, e.g., Johnson & Johnson Cordis Corporation, that make drug coated stents that keep arteries from clogging by releasing medication. Examples of M/N tools may be self-assembling, e.g., Angstrom Medica altered calcium and phosphate molecules that self-assemble to create nanostructured synthetic bone.

Another tool example is S. Stupp project at Northwestern University, which provides long complex molecules with hydrophobic tails and hydrophilic heads; these molecules self-assemble to form cylindrical structures that can be applied to making artificial bone. Another example of M/N tools 405 is Son Binh Nguyen use of nanoparticles for small molecule chemotherapy, in which engineered hydrophobic cyclic peptides attaches to targeted molecules and subsequently chemically react with molecule, breaking it into pieces.

Figure 4c shows components of therapeutic reservoirs 109. Release of therapies is dictated or controlled by systems-biology platform 104 instructions or from external source through wireless communication unit 106; timing mechanisms or rate of release may be reconfigured by software, e.g., MicroCHIPS implantable bioMEMS for drug delivery, in which silicon reservoirs hold medications in solid, liquid, or gel form, or iMEDD "NanoPORE Membranes,"

silicon wafers that have channels or pores with dimensions on nanometer scale for drug release.

Pre-filled reservoirs 410 contain medication filled-in biosensor 100 before implantation in living system. Contents of pre-filled reservoirs 410 may be pharmaceuticals or biopharmaceuticals in active form for release directly to living system. Pre-filled reservoirs 410 may hold probes, amino acids, nucleotides, or building blocks for sensor manufacture 110 for making additional biosensors.

Precursors 411 may be biological and chemical precursors to therapeutic pharmaceuticals and biopharmaceuticals. Depending on instructions from systems-biology platform 104, therapeutic precursors may be released, or therapeutic manufacture 108 may process into active pharmaceuticals or biopharmaceuticals.

Therapeutic storage 412 may store excess medication produced by therapeutic manufacture 108. Application of storing medication rather than manufacturing as needed if large doses, i.e., that cannot be made fast enough by therapeutic manufacture 108, are needed at time intervals.

Figure 4d shows basic components or interactions of sensor manufacture 110. Systems-biology platform 104 sends software instructions to sensor manufacture 110 to dictate manufacture, disassembly, activation, or deactivation of software-programmable biosensors. Once reconfigurable biosensors are programmed and produced, such components and sensor data signals are integrated, multiplexed, or processed

in combination into biosensor platform 102 for biological sensing.

Biosensor chip 421 acts as array or probe arranger 420  
5 attach probes onto array. Probe arranger 420 may attach probe  
for assaying, according to instructions by systems-biology  
platform 104. Method of attaching by probe arranger 420 can  
be printing method (e.g., placing probes on array with  
10 automated machinery). Probes may be attached through  
microspotting, in which automated microarray is produced by  
printing small quantities of pre-made biochemical substances  
onto solid surfaces.

Printing method may be ink-jet printing, e.g., GeSiM;  
15 non-contact method places probes on array in which probes are  
sprayed on surface using piezoelectric or other propulsion to  
transfer biochemical substances from nozzles to solid  
surfaces, or directly placed. This method allows *in situ*  
synthesis, advantageously synthesizing oligonucleotides on-  
20 the-fly directly on array surface. To change DNA that may be  
placed on array, systems-biology platform 104 provides probe  
arranger 420 list of sequences to synthesize.

Another example of probe arranger 420 is  
25 photolithography, e.g., Affymetrix GeneChips.  
Photolithography allows oligonucleotides to be built base-by-  
base (e.g, proteins build amino acid-by-amino acid) on array  
surface by repeated cycles of photodeprotection and  
nucleotide or amino acid addition. Like ink-jet printing,  
30 this process allows building of M/N arrays without pre-  
existing probes and can generate probes *in situ* on surface of  
biosensor chip 421.

Customizable microarray platform e.g., CombiMatrix, including semiconductor-based desktop microarray platform may fabricate custom oligonucleotide biochips. Microarrays with  
5 unique content are designed and fabricated on-the-fly using software driven process to generate reagents electrochemically. DNA oligonucleotides are synthesized *in situ* according to probe sequence designed. Probe arranger 420 may use cell-positioning chip, e.g., Aviva chip, to provide  
10 living whole-cell arrays.

Optionally soft lithography may use stamp to pattern surfaces of array, using patterned elastomer based on program instructions to define microfluidic networks on surface.  
15

Figure 5 shows DNA unit 500, representing organization of sensors in biosensor platform 102, such as RNA sensor 202, peptide or protein sensor 203, etc. DNA unit 500 may include DNA sensor 201, DNA therapeutic manufacture 501, DNA  
20 therapeutic reservoirs 502, or DNA reconfigurable biosensor 503 together in same physical structure, which lay in close proximity with each other. DNA therapeutic manufacture 501 is structure-specific category of therapeutic manufacture 108. DNA therapeutic reservoirs 502 and DNA reconfigurable  
25 biosensor 503 are structure-specific categories of therapeutic reservoirs 109 and sensor manufacturer 110 respectively.

Sequential steps begin with input introduction into DNA  
30 unit 500, specifically DNA sensor 201. Raw data is transferred to systems-biology platform 104, a remote source. Systems-biology platform 104 processes information,

outputting data and giving instructions to DNA therapeutic manufacture 501, DNA therapeutic reservoirs 502, and DNA reconfigurable biosensor 503. DNA therapeutic manufacture 501, DNA therapeutic reservoirs 502, and DNA reconfigurable biosensor 503 perform instructed tasks, with DNA sensor 201 monitoring respective progress.

DNA sensor 201 monitors or senses organism response to therapies dispensed by DNA therapeutic manufacture 501, DNA therapeutic reservoirs 502, or DNA reconfigurable biosensor 503. Proximity of DNA sensor 201, DNA therapeutic manufacture 501, DNA therapeutic reservoirs 502, or DNA reconfigurable biosensor 503 within same unit facilitates monitoring from DNA sensor 201.

Ongoing feedback is transmitted from DNA sensor 201 to DNA therapeutic manufacture 501, DNA therapeutic reservoirs 502, and DNA reconfigurable biosensor 503, while responding continually to DNA sensor 201 raw data, creating cyclic system of monitoring or responding.

Figure 6 flow chart shows automated or computer-assisted diagnosis of therapy recommendations or reports for target host, which is identified initially for possible diagnosis or treatment 601. To determine if host benefit from diagnosis or treatment, host undergoes preliminary screening 602. Preliminary screening may be implemented through software form; host undergoes preliminary modeling 603.

Modeling or simulation is used to model appropriate components or characteristics of device. After preliminary modeling 603, behavior of model is verified for accuracy 604.

If behavior of model is not ok, biosensor 100 is modified 605, and preliminary modelling 603 is repeated. If behavior of model is ok 604, biosensor 100 is configured 606. Reconfigurable biosensor is made or programmed according to  
5 such model.

Reconfigurable biosensor may be verified to comply or adhere to FDA regulations 607. If biosensor does not comply or adhere, it is modified 608 and configuration 606 or  
10 verification of adherence to FDA regulations 607 is repeated. If biosensor does comply or adhere to FDA regulations, it is implanted or attached to host 609.

Biosensor is initialized to allow sensor or detection  
15 activity *in vivo* 610. Sensing or software is executed 611. Initialization of biosensor and execution of sensing or software may operate in sequential order or in parallel. Once biosensor and software is initialized, initial *in vivo* sensing begins 612. Sensor data is then used for *in vivo*  
20 modeling 613 via systems-biology platform 104. After *in vivo* modeling 613, biosensor 100 generates diagnosis or therapy recommendation 614.

Therapy recommendations may result in commands to  
25 therapeutic unit 615 for therapeutic manufacturing or dispensing. Ongoing feedback between initial *in vivo* sensing 612, diagnosis or therapy recommendations 614, or commands to therapeutic unit creates an automated sensing, modeling, and treatment cycle.  
30

Foregoing descriptions of specific embodiments of the invention have been presented for purposes of illustration

and description. They are not intended to be exhaustive or to limit the invention to precise forms disclosed. Modifications and variations are possible in light of above teaching.

5

Embodiments were chosen or described in order to explain principles and application of the invention, thereby enabling others skilled in the art to utilize the invention in various embodiments or modifications according to particular purpose contemplated. Scope of the invention is intended to be defined by claims appended hereto and equivalents.

10



CLAIMS:

1. A computer-based method for predicting therapeutic strategy comprising the steps of:
  - 5 representing one or more biological processes and molecular interactions as mathematical models;  
accessing a systems biology platform via a network to said modeled biological processes and molecular interactions in a biological system, wherein the systems
  - 10 biology platform comprises genomics database, transcriptomics database, proteomics database, metabolomics database, pharmacogenomics database, pharmacokinetics database, disease database or drug database;  
running a set of simulations based on the data in the
  - 15 models and administration of a drug; and  
generating a simulation report for predicting outcome of the interactions with the drug, thereby aiding in target discovery and validation.
- 20 2. The method of claim 1 further comprising:  
simulating molecular interactions of biological processes in a metabolic, immune or genetic disorder.
3. The method of claim 1 further comprising:
  - 25 predicting one or more kinetic interactions between a drug and biological molecules in a biological system.
4. The method of claim 3 further comprising:  
analyzing the kinetic interactions between the drug
- 30 and biological molecules in the biological system.

5. The method of claim 1 further comprising:  
predicting differential diagnosis of a metabolic,  
immune or genetic disorder in an identified sub-group of  
biological hosts.

5

6. The method of claim 1 further comprising:  
predicting adverse reaction to a drug in a biological  
system.

10 7. The method of claim 1, wherein the mathematical model  
and simulation applies to analysis of drug toxicity in a  
biological system.

15 8. The method of claim 1, wherein the generated  
simulation outcome is visually displayed.

9. The method of claim 1, wherein the generated  
simulation outcome is stored in a data storage retrievable  
through the network.

20

10. A computer-based method to assist in medical decision  
making comprising the steps of:

accessing electronically a storage via a network,  
wherein the storage comprises genomics database, proteomics  
25 database, pharmacogenomics database, metabolomics database,  
pharmacokinetics database, transcriptomics database,  
disease database or drug database; and

implementing knowledge-based techniques in a systems  
biology platform to provide therapy, diagnosis or other  
30 automated feedback.

11. The method of claim 10 further comprising:  
using mathematical modeling and simulation of  
biological processes in a systems biology platform for  
differential diagnosis of a metabolic, immune or genetic  
5 disorder in a biological system.
12. The method of claim 10 further comprising:  
predicting one or more kinetic interactions between a  
drug and biological molecules in a biological system.
- 10 13. The method of claim 12 further comprising:  
analyzing the kinetic interactions between the drug  
and biological molecules in the biological system.
- 15 14. The method of claim 10 further comprising:  
predicting adverse reaction to a drug in a biological  
system.
- 20 15. The method of claim 10 further comprising:  
analyzing drug toxicity in a biological system.
- 25 16. A system for computational modeling of disease in a  
biological host comprising:  
a processor; and  
a memory;  
wherein the processor represents mathematically via a  
network genomic, transcriptomic, metabolomic,  
pharmacogenomic or pharmacokinetic data, runs one or more  
simulations to analyze biological interactions and predicts  
30 the effect of a drug on disease.

17. The system of claim 16, wherein the biological interaction further comprises molecular interactions of biological processes in a metabolic, immune or genetic disorder in a biological system.

5

18. The system of claim 16, wherein the mathematical representation further comprises analysis of drug toxicity in a biological system.

10 19. The system of claim 16 further comprising:  
prediction of kinetic interactions between a drug and biological molecules in a biological system.

15 20. The system of claim 19 further comprising:  
analysis of the kinetic interactions between the drug and biological molecules in the biological system.

20 21. The system of claim 16 further comprising:  
prediction of adverse reaction to a drug in a biological system.

22. The system of claim 16 further comprising:  
analysis of drug toxicity in a biological system.

25 23. The system of claim 16 further comprising:  
an electronic data storage to store and access simulation data.

30 24. Implantable network-biosensor comprising:  
a sensor unit for receiving a multi-sensor signal from a biosensor platform for detecting a biological material of a host;

a controller for processing a systems-biology platform for verifying or modifying a simulation model associated with the biological material; and

5 a therapeutic unit for releasing or dispensing a therapeutic material from a reservoir in or onto the host, whereby the sensor unit may automatically detect an effect of the therapeutic material on the host.

25. The network biosensor of claim 24 wherein:

10 the sensor unit is configurable or programmable for detecting multi-sensor signaling, thereby enabling the biosensor platform to access one or more sensor signals from a DNA or RNA sensor, a peptide or protein sensor, an antibody or antigen sensor, a vector or virus-vector  
15 sensor, a lipid or fatty-acid sensor, or an inorganic-ion or electrochemical sensor.

26. The network biosensor of claim 24 wherein:

20 the sensor unit is configurable or programmable for detecting multi-sensor signaling, thereby enabling the biosensor platform to access one or more sensor signals from a tissue-factor sensor, a steroid sensor, a neurotransmitter sensor, a pH sensor, a free-radical  
25 sensor, a carbohydrate sensor, a neural sensor, a chemical sensor, a small-molecule sensor, an exon sensor, a metabolites sensor, an intermediate sensor, a chromosome sensor, or a cell sensor.

27. The network biosensor of claim 24 wherein:

30 the sensor unit comprises a positioning chip for immobilizing or positioning the biological material for sensing thereof.

28. The network biosensor of claim 24 wherein:

the sensor unit receives another multi-sensor signal from another biosensor platform for detecting another biological material of the same or another host.

5

29. The network biosensor of claim 24 wherein:

the controller is configurable or programmable for processing multiple simulation applications, thereby enabling the systems-biology platform to access one or more simulation models comprising a genomics model, a proteomics model, a computational chemistry model, a pharmacogenomics model, a computational biology model, a computational biophysics model, a computational cell behavior model, a pharmacokinetics model, a metabolomics model, or a transcriptomics model.

10

15

30. The network biosensor of claim 24 wherein:

the controller is configurable or programmable for processing multiple simulation data, thereby enabling the systems-biology platform to access one or more simulation data comprising a genetic-disorder or mutation data, an infectious disease or infection data, an immunity-disease data, a single-organ or cell-type autoimmune disease data, or a neoplasia data.

20

25

31. The network biosensor of claim 24 further comprising:

a therapeutic unit for releasing or dispensing a therapeutic material from a reservoir in or onto the host, whereby the sensor unit may automatically detect an effect of the therapeutic material on the host.

30

32. The network biosensor of claim 24 wherein:

the therapeutic unit is configurable or programmable for releasing or dispensing the therapeutic material alternatively from manufacture means, thereby enabling the  
5 systems-biology platform to instruct the therapeutic unit configurably or programmably using one or more manufacture-means components comprising pharmaceuticals, biopharmaceuticals, reconfigurable biocatalytic chips, tissue scaffolds, or micro or nano-array or electro-  
10 mechanical tools.

33. The network biosensor of claim 24 wherein:

the controller processes the systems-biology platform adaptively for generating a diagnostic or therapeutic  
15 signal or report, whereby the systems-biology platform may access one or more simulation applications comprising a neural or learning network, a statistical or probabilistic expert, fuzzy-logic or knowledge-based system, an artificial intelligence or decision-making inference-engine  
20 or program, or a supervised or unsupervised Bayesian or Markovian analysis, clustering, criterion or classification program.

**Application No:** GB0722767.1

**Examiner:** Dr Patrick Purcell

**Claims searched:** 24-33

**Date of search:** 6 December 2007

## Patents Act 1977: Search Report under Section 17

### Documents considered to be relevant:

Category	Relevant to claims	Identity of document and passage or figure of particular relevance
Y	24-26, 29-33	US6178349 A (KIEVAL) see whole document, esp. column 3, line 65-column 5, line 36, column 8, lines 4-19, column 11, lines 29-44
Y	24-26, 29-33	WO03/043684 A (GLAXO GROUP LIMITED) see whole document, esp. page 3, line 2-page 7, line 15, page 8, lines 18-32, page 19, lines 4-20

### Categories:

X	Document indicating lack of novelty or inventive step	A	Document indicating technological background and/or state of the art.
Y	Document indicating lack of inventive step if combined with one or more other documents of same category.	P	Document published on or after the declared priority date but before the filing date of this invention.
&	Member of the same patent family	E	Patent document published on or after, but with priority date earlier than, the filing date of this application.

### Field of Search:

Search of GB, EP, WO & US patent documents classified in the following areas of the UKC<sup>x</sup> :

Worldwide search of patent documents classified in the following areas of the IPC

The following online and other databases have been used in the preparation of this search report

ONLINE: EPODOC, WPI

### International Classification:

Subclass	Subgroup	Valid From
G06F	0019/00	01/01/2006
A61B	0005/00	01/01/2006
G01N	0033/48	01/01/2006